

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt SEQUENCE LISTING

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<110> Genentech, Inc.
Ashkenazi, Avi
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          Eaton, Dan L.
          Ferrara, Napoleone
Filvaroff, Ellen
          Fong, Sherman
Gao, Wei-Qiang
          Gerber, Hanspeter
          Gerritsen, Mary E.
          Goddard, A.
Godowski, Paul J.
Grimaldi, Christopher J.
          Gurney, Austin L.
          Hillan, Kenneth, J.
          Kljavin, Ivar J.
Mather, Jennie P.
          Pan, James
          Paoni, Nicholas F.
          Roy, Margaret Ann
          Stewart, Timothy A.
          Tumas, Daniel
          Williams, P. Mickey Wood, William, I.
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt ctcaattagt cagcaaccca gtttt

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2	51. 1
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Pro Ala Ser Tyr Arg Leu Trp Gly Ala Pro Leu Gln Pro Thr Leu Gly 50 60
Val Val Pro Gln Ala Ser Val Pro Leu Leu Thr Asp Leu Ala Gln Trp 65 70 75 80
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Tyr Val Cys Thr Pro Val Pro His Pro Asp Pro Pro Met Ala Leu Ser
Arg Thr Pro Thr Arg Gln Ile Ser Ser Ser Asp Thr Asp Pro Pro Ala
115 120 125
Asp Gly Pro Ser Asn Pro Leu Cys Cys Cys Phe His Gly Pro Ala Phe
Ser Thr Leu Asn Pro Val Leu Arg His Leu Phe Pro Gln Glu Ala Phe
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Arg Thr Ser Gly Lys His Val Gln Val Thr Gly Arg Arg Ile Ser Ala 50 60
Thr Ala Glu Asp Gly Asn Lys Phe Ala Lys Leu Ile Val Glu Thr Asp 65 70 75 80
Thr Phe Gly Ser Arg Val Arg Ile Lys Gly Ala Glu Ser Glu Lys Tyr
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Ile Cys Met Asn Lys Arg Gly Lys Leu Ile Gly Lys Pro Ser Gly Lys 100 105 110
Ser Lys Asp Cys Val Phe Thr Glu Ile Val Leu Glu Asn Asn Tyr Thr
115
Ala Phe Gln Asn Ala Arg His Glu Gly Trp Phe Met Ala Phe Thr Arg
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Gln Gly Arg Pro Arg Gln Ala Ser Arg Ser Arg Gln Asn Gln Arg Glu
145 150 155 160
Ala His Phe Ile Lys Arg Leu Tyr Gln Gly Gln Leu Pro Phe Pro Asn
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly Ile Pro Glu Gly 50 55 60
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Pro Ala Glu Leu His Asn Val Gln Ser Val His Thr Val Tyr Leu Tyr
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Gly Asn Gln Leu Asp Glu Phe Pro Met Asn Leu Pro Lys Asn Val Arg
Val Leu His Leu Gln Glu Asn Asn Ile Gln Thr Ile Ser Arg Ala Ala
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Leu Ala Gln Leu Leu Lys Leu Glu Glu Leu His Leu Asp Asp Asn Ser
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Ile Ser Thr Val Gly Val Glu Asp Gly Ala Phe Arg Glu Ala Ile Ser
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Leu Lys Leu Leu Phe Leu Ser Lys Asn His Leu Ser Ser Val Pro Val
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Ala Val Ile Ser Asp Met Ala Phe Gln Asn Leu Thr Ser Leu Glu Arg
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Leu Ile Val Asp Gly Asn Leu Leu Thr Asn Lys Gly Ile Ala Glu Gly 210 215 220
Thr Phe Ser His Leu Thr Lys Leu Lys Glu Phe Ser Ile Val Arg Asn 230 235 240
225
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Ser Asn Leu Arg Lys Leu Glu Arg Leu Asp Ile Ser Asn Asn Gln Leu 275 280 285 Arg Met Leu Thr Gln Gly Val Phe Asp Asn Leu Ser Asn Leu Lys Gln 290 295 300 Leu Thr Ala Arg Asn Asn Pro Trp Phe Cys Asp Cys Ser Ile Lys Trp 305 310 315 Val Thr Glu Trp Leu Lys Tyr Ile Pro Ser Ser Leu Asn Val Arg Gly 325 330 335 Phe Met Cys Gln Gly Pro Glu Gln Val Arg Gly Met Ala Val Arg Glu 340 345 350 Leu Asn Met Asn Leu Leu Ser Cys Pro Thr Thr Pro Gly Leu Pro 355 360 365 Leu Phe Thr Pro Ala Pro Ser Thr Ala Ser Pro Thr Thr Gln Pro Pro 370 380 Thr Leu Ser Ile Pro Asn Pro Ser Arg Ser Tyr Thr Pro Pro Thr Pro 385 390 395 Thr Thr Ser Lys Leu Pro Thr Ile Pro Asp Trp Asp Gly Arg Glu Arg 405 410 415 Val Thr Pro Pro Ile Ser Glu Arg Ile Gln Leu Ser Ile His Phe Val 420 425 430 Asn Asp Thr Ser Ile Gln Val Ser Trp Leu Ser Leu Phe Thr Val Met 435 440 445 Ala Tyr Lys Leu Thr Trp Val Lys Met Gly His Ser Leu Val Gly Gly 450 460 Ile Val Gln Glu Arg Ile Val Ser Gly Glu Lys Gln His Leu Ser Leu 465 470 475 480 Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg Ile Cys Leu Val Pro Leu 485 490 495 Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp Thr Ile Cys Ser Glu Ala 500 505 Thr Thr His Ala Ser Tyr Leu Asn Asn Gly Ser Asn Thr Ala Ser Ser 515 520 525 His Glu Gln Thr Thr Ser His Ser Met Gly Ser Pro Phe Leu Leu Ala 530 540 Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser 545 550 555 560 Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys 565 570 575 Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp Asp Tyr Cys Glu Ala Gly 580 585 Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln 595 600 605

~ 60 . The second results of ~ 10 . We have the second results are ~ 10 .	
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Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His 625 630 635 640	•
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Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys 100 105 110

Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 115 120 125

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Met Cys Ser Thr Leu Glu His Asn Cys Ala His Phe Cys Ile Asn Ile 245 250 255

Pro Gly Ser Tyr Val Cys Arg Cys Lys Gln Gly Tyr Ile Leu Asn Ser 260 265 270

Asp Gln Thr Thr Cys Arg Ile Gln Asp Leu Cys Ala Met Glu Asp His 275 280 285

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Ala Ala Ile Val Phe Thr Asp Gly Arg Ala Gln Asp Asp Val Ser Glu 725 730 735

Trp Ala Ser Lys Ala Lys Ala Asn Gly Ile Thr Met Tyr Ala Val Gly
740 745 750

Val Gly Lys Ala Ile Glu Glu Glu Leu Gln Glu Ile Ala Ser Glu Pro 755 760 765

Thr Asn Lys His Leu Phe Tyr Ala Glu Asp Phe Ser Thr Met Asp Glu 770 775 780

Ile Ser Glu Lys Leu Lys Lys Gly Ile Cys Glu Ala Leu Glu Asp Ser 785 790 795 800

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Gln Pro Thr Glu Ser Glu Pro Val Thr Ile Asn Ile Gln Asp Leu Leu 820 825 830

Ser Cys Ser Asn Phe Ala Val Gln His Arg Tyr Leu Phe Glu Glu Asp 835 840 845

Asn Leu Leu Arg Ser Thr Gln Lys Leu Ser His Ser Thr Lys Pro Ser 850 855 860

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Ile Met Phe Gln Asn Leu Ala Asn Glu Glu Val Arg Lys Leu Thr Gln 885 890 895

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<400> 35

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<220>
 <223> Description of Artificial Sequence: Synthetic
         oligonucleotide probe
<400> 36
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<211> 45
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 <213> Artificial Sequence
<220>
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         oligonucleotide probe
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<212> DNA
<213> Homo sapiens
<400> 38
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ccttggtcta ctccatgccc tcccggaacc tgtccctgcg gctggagggt ctccaggaga 480
aagactctgg cccctacagc tgctccgtga atgtgcaaga caaacaaggc aaatctaggg 540
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attgggagga gcctccaccc acccctgact cctccttatg aagccagctg ctgaaattag 1560
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<213> Homo sapiens
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Leu His Leu Pro Ala Asn Arg Leu Gln Ala Val Glu Gly Glu Val
35 40 45
Val Leu Pro Ala Trp Tyr Thr Leu His Gly Glu Val Ser Ser Gln
50 55 60
Pro Trp Glu Val Pro Phe Val Met Trp Phe Phe Lys Gln Lys Glu Lys 65 70 75 80
Glu Asp Gln Val Leu Ser Tyr Ile Asn Gly Val Thr Thr Ser Lys Pro
85 90 95
Gly Val Ser Leu Val Tyr Ser Met Pro Ser Arg Asn Leu Ser Leu Arg
100 105 110
Leu Glu Gly Leu Gln Glu Lys Asp Ser Gly Pro Tyr Ser Cys Ser Val
115 120 125
Asn Val Gln Asp Lys Gln Gly Lys Ser Arg Gly His Ser Ile Lys Thr
130 135 140
Leu Glu Leu Asn Val Leu Val Pro Pro Ala Pro Pro Ser Cys Arg Leu
145 150 155 160
Gln Gly Val Pro His Val Gly Ala Asn Val Thr Leu Ser Cys Gln Ser
165 170 175
Pro Arg Ser Lys Pro Ala Val Gln Tyr Gln Trp Asp Arg Gln Leu Pro
180 185 190
Ser Phe Gln Thr Phe Phe Ala Pro Ala Leu Asp Val Ile Arg Gly Ser
195 200 205
Leu Ser Leu Thr Asn Leu Ser Ser Ser Met Ala Gly Val Tyr Val Cys
210 215 220
Lys Ala His Asn Glu Val Gly Thr Ala Gln Cys Asn Val Thr Leu Glu
225 230 235 240
Val Ser Thr Gly Pro Gly Ala Ala Val Val Ala Gly Ala Val Val Gly 245 250 255
Thr Leu Val Gly Leu Gly Leu Leu Ala Gly Leu Val Leu Leu Tyr His 260 265 270
Arg Arg Gly Lys Ala Leu Glu Glu Pro Ala Asn Asp Ile Lys Glu Asp 275 280 285
Ala Ile Ala Pro Arg Thr Leu Pro Trp Pro Lys Ser Ser Asp Thr Ile
290 295 300
                                             Page 22
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Ser Lys Asn Gly Thr Leu Ser Ser Val Thr Ser Ala Arg Ala Leu Arg 305 310 Pro Pro His Gly Pro Pro Arg Pro Gly Ala Leu Thr Pro Thr Pro Ser Leu Ser Ser Gln Ala Leu Pro Ser Pro Arg Leu Pro Thr Thr Asp Gly 340 Ala His Pro Gln Pro Ile Ser Pro Ile Pro Gly Gly Val Ser Ser Ser Gly Leu Ser Arg Met Gly Ala Val Pro Val Met Val Pro Ala Gln Ser Gln Ala Gly Ser Leu Val 385 <210> 40 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 40 22 agggtctcca ggagaaagac tc <210> 41 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 41 24 attgtgggcc ttgcagacat agac <210> 42 <211> 50 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic oligonucleotide probe 50 ggccacagca tcaaaacctt agaactcaat gtactggttc ctccagctcc <210> 43 <211> 18 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

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<210> 46 <211> 24 <212> DNA <213> Artificial Sequence	,
<220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe	•
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<210> 47 <211> 45 <212> DNA <213> Artificial Sequence	
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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 caaactgcca titagataat gtctgtatag ctgcaaatat taataaaact ttaacaaaaa 420
tcagatccat aaaagaacct gtggctttgc tacaagaagt ctatagaaat tctgtgacag 480
atcittcacc aacagatata attacatata tagaaatait agctgaatca tcttcattac 540
taggttacaa gaacaacact atctcagcca aggacaccct ttctaactca actcttactg 600
aatitgtaaa aaccgtgaat aattttgttc aaagggatac atttgtagtt tgggacaagt 660
tatctgtgaa tcataggaga acacatctta caaaactcat gcacactgtt gaacaagcta 720 ctttaaggat atcccagagc ttccaaaaga ccacagggt tgatacaaat tcaacggata 780 tagctctcaa agttttcttt tttgattcat ataacatgaa acatattcat cctcatatga 840 atatggatgg aggtgcattt ttatattata agagtattgg tcctttgctt tcatcatctg 900 gcaatgttgc agttgcattt ttatattata agagtattgg tcctttgctt tcatcatctg 960 acaacttctt attgaaacct caaaattatg ataattctga agaggaggaa agagtcatat 1020 cttcagtaat ttcagtctca atgagctcaa acccaccac attatataaa cttgaaaaaa 1080
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 tgtataaaaa tgactcatca aattatccaa ttattaacta ctagacaaaa agtattttaa 2220
 atcagttttt ctgtttatgc tataggaact gtagataata aggtaaaatt atgtatcata 2280
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Ser Tyr Thr Gln Asn Cys Thr Lys Thr Pro Cys Leu Pro Asn Ala Lys
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Cys Glu Ile Arg Asn Gly Ile Glu Ala Cys Tyr Cys Asn Met Gly Phe 35 40 45
Ser Gly Asn Gly Val Thr Ile Cys Glu Asp Asp Asn Glu Cys Gly Asn 50 60
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Leu Thr Gln Ser Cys Gly Glu Asn Ala Asn Cys Thr Asn Thr Glu Gly

Ser Tyr Tyr Cys Met Cys Val Pro Gly Phe Arg Ser Ser Ser Asn Gln
85 90 95

65

Asp Arg Phe Ile Thr Asn Asp Gly Thr Val Cys Ile Glu Asn Val Asn 100 105 110

Ala Asn Cys His Leu Asp Asn Val Cys Ile Ala Ala Asn Ile Asn Lys 115 120 125

Thr Leu Thr Lys Ile Arg Ser Ile Lys Glu Pro Val Ala Leu Leu Gln 130 135 140

Glu Val Tyr Arg Asn Ser Val Thr Asp Leu Ser Pro Thr Asp Ile Ile 145 150 155 160

Thr Tyr Ile Glu Ile Leu Ala Glu Ser Ser Ser Leu Leu Gly Tyr Lys 165 170 175

Asn Asn Thr Ile Ser Ala Lys Asp Thr Leu Ser Asn Ser Thr Leu Thr 180 185 190

Glu Phe Val Lys Thr Val Asn Asn Phe Val Gln Arg Asp Thr Phe Val 195 200 205

val Trp Asp Lys Leu Ser Val Asn His Arg Arg Thr His Leu Thr Lys 210 220

Leu Met His Thr Val Glu Gln Ala Thr Leu Arg Ile Ser Gln Ser Phe 225 230 235 240

Gln Lys Thr Thr Glu Phe Asp Thr Asn Ser Thr Asp Ile Ala Leu Lys 245 250 255

Val Phe Phe Phe Asp Ser Tyr Asn Met Lys His Ile His Pro His Met 260 265 270

Asn Met Asp Gly Asp Tyr Ile Asn Ile Phe Pro Lys Arg Lys Ala Ala 275 280 285

Tyr Asp Ser Asn Gly Asn Val Ala Val Ala Phe Leu Tyr Tyr Lys Ser 290 295 300

Ile Gly Pro Leu Leu Ser Ser Ser Asp Asn Phe Leu Leu Lys Pro Gln 305 310 315 320

Asn Tyr Asp Asn Ser Glu Glu Glu Glu Arg Val Ile Ser Ser Val Ile 325 330 335

Ser Val Ser Met Ser Ser Asn Pro Pro Thr Leu Tyr Glu Leu Glu Lys 340 345 350

Ile Thr Phe Thr Leu Ser His Arg Lys Val Thr Asp Arg Tyr Arg Ser 355 360 365

Leu Cys Ala Phe Trp Asn Tyr Ser Pro Asp Thr Met Asn Gly Ser Trp 370 375 380

Ser Ser Glu Gly Cys Glu Leu Thr Tyr Ser Asn Glu Thr His Thr Ser 385 390 395 400

Cys Arg Cys Asn His Leu Thr His Phe Ala Ile Leu Met Ser Ser Gly Page 26

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1,txt
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                   405
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Leu Gly Ile Ile Ser Leu Ile Cys Leu Ala Ile Cys Ile Phe Thr
435 440 445
Phe Trp Phe Phe Ser Glu Ile Gln Ser Thr Arg Thr Thr Ile His Lys 450 455 460
Asn Leu Cys Cys Ser Leu Phe Leu Ala Glu Leu Val Phe Leu Val Gly
465 470 475 480
Ile Asn Thr Asn Thr Asn Lys Leu Phe Cys Ser Ile Ile Ala Gly Leu
485 490 495
Leu His Tyr Phe Phe Leu Ala Ala Phe Ala Trp Met Cys Ile Glu Gly 500 510
Ile His Leu Tyr Leu Ile Val Val Gly Val Ile Tyr Asn Lys Gly Phe 515 520 525
Leu His Lys Asn Phe Tyr Ile Phe Gly Tyr Leu Ser Pro Ala Val 530 540
Val Gly Phe Ser Ala Ala Leu Gly Tyr Arg Tyr Tyr Gly Thr Thr Lys
545 550 555 560
Val Cys Trp Leu Ser Thr Glu Asn Asn Phe Ile Trp Ser Phe Ile Gly
565 570 575
Pro Ala Cys Leu Ile Ile Leu Val Asn Leu Leu Ala Phe Gly Val Ile
580 585 590
Ile Tyr Lys Val Phe Arg His Thr Ala Gly Leu Lys Pro Glu Val Ser
595 600 605
Cys Phe Glu Asn Ile Arg Ser Cys Ala Arg Gly Ala Leu Ala Leu Leu 610 615 620
Phe Leu Leu Gly Thr Thr Trp Ile Phe Gly Val Leu His Val Val His 625 630 635 640
Ala Ser Val Val Thr Ala Tyr Leu Phe Thr Val Ser Asn Ala Phe Gln
Gly Met Phe Ile Phe Leu Phe Leu Cys Val Leu Ser Arg Lys Ile Gln 660 670
Glu Glu Tyr Tyr Arg Leu Phe Lys Asn Val Pro Cys Cys Phe Gly Cys 675 680 685
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Leu Arg
690
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<213> Homo sapiens
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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gtattggtcc ctttgctttc atcatctgac aacttcttat tgaaacctca aaattatgat 180
aattctgaag aggaggaaag agtcatatct tcagtaattt cagtctcaat gagctcaaac 240 ccacccacat tatatgaact tgaaaaaata acatttacat taagtcatcg aaaggtcaca 300 gataggtata ggagtctatg tggcattttg gaatactcac ctgataccat gaatggcagc 360 tggtcttcag agggctgtag gctgacatac tcaaatgaga cccacacctc atgccgctgt 420 aatcacctga cacattttgc aattttgatg tcctctggtc cttccattgg tattaaagat 480
tataatatic ttacaaggat cactcaacta ggaataatta tttcactgat ttgtcttgcc 540
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<210> 51
<211> 20
<212> DNA
<213> Artificial Sequence
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       oligonucleotide probe
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 52
                                                                                    18
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<210> 53
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 53
                                                                                    22
cacctgatac catgaatggc ag
<210> 54
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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                                                                                      18
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<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
        oligonucleotide probe
<400> 56
                                                                                      23
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<210> 57
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<212> DNA
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        oligonucleotide probe
                                                                                      50
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<211> 2137
<212> DNA
<213> Homo sapiens
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                                                  Page 29
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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gcťgcačtgt cťcagttčtg cťťgaatacc tccaťcgatg gggaactcac tťcctttgga 1440
āaaāttetīa tgteāagetīg aaaīteteta attttītetē ātēaettece caggageāge 1500
cagaagacag gcagtagtti taatttcagg aacaggtgat ccactctgta aaacagcagg 1560
taaatttcac tcaaccccat gtgggaattg atctatatct ctacttccag ggaccatttg 1620
cccttcccaa atccctccag gccagaactg actggagcag gcatggccca ccaggcttca
ggagtagggg aagcctggag ccccactcca gccctgggac aacttgagaa ttccccctga
ggccagttct gtcatggatg ctgtcctgag aataacttgc tgtcccggtg tcacctgctt
1980
                                                                         2040
īgtacatctc ttatīttctt ācattattta tgcccccaaa ttātātttat gtatgtāagt
                                                                         2100
gaggtttgtt ttgtatatta aaatggagtt tgtttgt
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<211> 216
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His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg His Leu Tyr 35 40 45
Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu Arg Ile Arg Ala 50 55 60
Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser Ala His Ser Leu Leu 65 70 75 80
Glu Ile Lys Ala Val Ala Leu Arg Thr Val Ala Ile Lys Gly Val His
85 90 95
Ser Val Arg Tyr Leu Cys Met Gly Ala Asp Gly Lys Met Gln Gly Leu 100 \hspace{1cm} 105 \hspace{1cm} 110
Leu Gln Tyr Ser Glu Glu Asp Cys Ala Phe Glu Glu Glu Ile Arg Pro
115 120 125
Asp Gly Tyr Asn Val Tyr Arg Ser Glu Lys His Arg Leu Pro Val Ser
130 135 140
Leu Ser Ser Ala Lys Gln Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu
145 150 155 160
Pro Leu Ser His Phe Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro
Glu Asp Leu Arg Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu 180 185 190
Glu Thr Asp Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala
195 200 205
Val Arg Ser Pro Ser Phe Glu Lys
210 215
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

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         oligonucleotide probe
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                                                                                                            26
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<210> 61
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<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
         oligonucleotide probe
                                                                                                            42
gcctcccggt ctccctgagc agtgccaaac agcggcagtg ta
<210> 62
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
         oligonucleotide probe
<400> 62
                                                                                                            22
ccagtccggt gacaagccca aa
<210> 63
<211> 1295
<212> DNA
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Glu Ala Ser Thr Val Asp Cys Asn Asp Leu Gly Leu Leu Thr Phe Pro
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Ala Arg Leu Pro Ala Asn Thr Gln Ile Leu Leu Gln Thr Asn Asn 65 70 75 80
Ile Ala Lys Ile Glu Tyr Ser Thr Asp Phe Pro Val Asn Leu Thr Gly
85 90 95
Leu Asp Leu Ser Gln Asn Asn Leu Ser Ser Val Thr Asn Ile Asn Val
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Lys Lys Met Pro Gln Leu Leu Ser Val Tyr Leu Glu Glu Asn Lys Leu
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Thr Glu Leu Pro Glu Lys Cys Leu Ser Glu Leu Ser Asn Leu Gln Glu
130 135 140
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Leu Tyr Ile Asn His Asn Leu Leu Ser Thr Ile Ser Pro Gly Ala Phe Ile Gly Leu His Asn Leu Leu Arg Leu His Leu Asn Ser Asn Arg 165 Gln Met Ile Asn Ser Lys Trp Phe Asp Ala Leu Pro Asn Leu Glu Ile 180 185 190 Leu Met Ile Gly Glu Asn Pro Ile Ile Arg Ile Lys Asp Met Asn Phe 195 200 205 Lys Pro Leu Ile Asn Leu Arg Ser Leu Val Ile Ala Gly Ile Asn Leu 210 220 Glu Ile Pro Asp Asn Ala Leu Val Gly Leu Glu Asn Leu Glu Ser 230 235 240 Ile Ser Phe Tyr Asp Asn Arg Leu Ile Lys Val Pro His Val Ala Leu 245 250 255 Gln Lys Val Val Asn Leu Lys Phe Leu Asp Leu Asn Lys Asn Pro Ile 260 265 270 Asn Arg Ile Arg Arg Gly Asp Phe Ser Asn Met Leu His Leu Lys Glu 275 280 285 Leu Gly Ile Asn Asn Met Pro Glu Leu Ile Ser Ile Asp Ser Leu Ala 290 295 300 Val Asp Asn Leu Pro Asp Leu Arg Lys Ile Glu Ala Thr Asn Asn Pro 305 310 315 320 Arg Leu Ser Tyr Ile His Pro Asn Ala Phe Phe Arg Leu Pro Lys Leu 325 330 335 Glu Ser Leu Met Leu Asn Ser Asn Ala Leu Ser Ala Leu Tyr His Gly 340 350 Thr Ile Glu Ser Leu Pro Asn Leu Lys Glu Ile Ser Ile His Ser Asn 355 360 365 Pro Ile Arg Cys Asp Cys Val Ile Arg Trp Met Asn Met Asn Lys Thr 370 375 380 Asn Ile Arg Phe Met Glu Pro Asp Ser Leu Phe Cys Val Asp Pro Pro 385 390 395 400 Glu Phe Gln Gly Gln Asn Val Arg Gln Val His Phe Arg Asp Met Met 405 410 415 Glu Ile Cys Leu Pro Leu Ile Ala Pro Glu Ser Phe Pro Ser Asn Leu 420 425 430 Asn Val Glu Ala Gly Ser Tyr Val Ser Phe His Cys Arg Ala Thr Ala 435 440 445 Glu Pro Gln Pro Glu Ile Tyr Trp Ile Thr Pro Ser Gly Gln Lys Leu 450 455 460 Leu Pro Asn Thr Leu Thr Asp Lys Phe Tyr Val His Ser Glu Gly Thr 465 470 475 480

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Ile Arg Asp Ile Gln Ala Asn Ser Val Leu Val Ser Trp Lys Ala Ser
530 540
Ser Lys Ile Leu Lys Ser Ser Val Lys Trp Thr Ala Phe Val Lys Thr 545 550 555 560
Glu Asn Ser His Ala Ala Gln Ser Ala Arg Ile Pro Ser Asp Val Lys
565 575
Val Tyr Asn Leu Thr His Leu Asn Pro Ser Thr Glu Tyr Lys Ile Cys
580 585 590
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val Thr Thr Lys Gly Leu His Pro Asp Gln Lys Glu Tyr Glu Lys Asn
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625 630 635 640
Gly Val Ile Cys Leu Ile Ser Cys Leu Ser Pro Glu Met Asn Cys Asp
645 650 655
Gly Gly His Ser Tyr Val Arg Asn Tyr Leu Gln Lys Pro Thr Phe Ala
660 670
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Glu Lys Ser Thr Ser Leu Lys Val Lys Ala Thr Val Ile Gly Leu Pro
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Thr Cys Ser Asn Ala Asn Leu Lys Glu Ile Pro Arg Asp Leu Pro Pro 50 60
Glu Thr Val Leu Leu Tyr Leu Asp Ser Asn Gln Ile Thr Ser Ile Pro
65 70 75 80
Asn Glu Ile Phe Lys Asp Leu His Gln Leu Arg Val Leu Asn Leu Ser
85 90 95
Lys Asn Gly Ile Glu Phe Ile Asp Glu His Ala Phe Lys Gly Val Ala
100 105 110
Glu Thr Leu Gln Thr Leu Asp Leu Ser Asp Asn Arg Ile Gln Ser Val
115 120 125
His Lys Asn Ala Phe Asn Asn Leu Lys Ala Arg Ala Arg Ile Ala Asn
130 135 140
Asn Pro Trp His Cys Asp Cys Thr Leu Gln Gln Val Leu Arg Ser Met
145 150 155 160
Ala Ser Asn His Glu Thr Ala His Asn Val Ile Cys Lys Thr Ser Val
165 170 175
Leu Asp Glu His Ala Gly Arg Pro Phe Leu Asn Ala Ala Asn Asp Ala
180 185 190
Asp Leu Cys Asn Leu Pro Lys Lys Thr Thr Asp Tyr Ala Met Leu Val
195 200 205
Thr Met Phe Gly Trp Phe Thr Met Val Ile Ser Tyr Val Val Tyr Tyr
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Ser Val Leu Ser Gly Ser Ala Thr Gly Cys Pro Pro Arg Cys Glu Cys Ser Ala Gln Asp Arg Ala Val Leu Cys His Arg Lys Cys Phe Val Ala 50 60 Val Pro Glu Gly Ile Pro Thr Glu Thr Arg Leu Leu Asp Leu Gly Lys 65 70 75 80 Asn Arg Ile Lys Thr Leu Asn Gln Asp Glu Phe Ala Ser Phe Pro His 85 90 95 Leu Glu Glu Leu Glu Leu Asn Glu Asn Ile Val Ser Ala Val Glu Pro 100 105 110 Gly Ala Phe Asn Asn Leu Phe Asn Leu Arg Thr Leu Gly Leu Arg Ser 115 120 125 Asn Arg Leu Lys Leu Ile Pro Leu Gly Val Phe Thr Gly Leu Ser Asn 130 135 140 Leu Thr Lys Gln Asp Ile Ser Glu Asn Lys Ile Val Ile Leu Leu Asp 145 150 155 160 Tyr Met Phe Gln Asp Leu Tyr Asn Leu Lys Ser Leu Glu Val Gly Asp 165 170 175 Asn Asp Leu Val Tyr Ile Ser His Arg Ala Phe Ser Gly Leu Asn Ser 180 185 190 Leu Glu Gln Leu Thr Leu Glu Lys Cys Asn Leu Thr Ser Ile Pro Thr 195 200 205 Glu Ala Leu Ser His Leu His Gly Leu Ile Val Leu Arg Leu Arg His 210 215 220 Leu Asn Ile Asn Ala Ile Arg Asp Tyr Ser Phe Lys Arg Leu Tyr Arg 225 230 235 240 Leu Lys Val Leu Glu Ile Ser His Trp Pro Tyr Leu Asp Thr Met Thr 245 250 255 Pro Asn Cys Leu Tyr Gly Leu Asn Leu Thr Ser Leu Ser Ile Thr His 260 265 270 Cys Asn Leu Thr Ala Val Pro Tyr Leu Ala Val Arg His Leu Val Tyr 275 280 285 Leu Arg Phe Leu Asn Leu Ser Tyr Asn Pro Ile Ser Thr Ile Glu Gly 290 295 300 Met Leu His Glu Leu Leu Arg Leu Gln Glu Ile Gln Leu Val Gly 310 315 320 Gly Gln Leu Ala Val Val Glu Pro Tyr Ala Phe Arg Gly Leu Asn Tyr 325 330 335 Leu Arg Val Leu Asn Val Ser Gly Asn Gln Leu Thr Thr Leu Glu Glu 340 345 350 Ser Val Phe His Ser Val Gly Asn Leu Glu Thr Leu Ile Leu Asp Ser 355 360 365

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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385 390 395 400
Phe Val Gln Gly Lys Glu Phe Lys Asp Phe Pro Asp Val Leu Leu Pro 405 410 415
Asn Tyr Phe Thr Cys Arg Arg Ala Arg Ile Arg Asp Arg Lys Ala Gln
420 425 430
Gln Val Phe Val Asp Glu Gly His Thr Val Gln Phe Val Cys Arg Ala
435 440 445
Asp Gly Asp Pro Pro Pro Ala Ile Leu Trp Leu Ser Pro Arg Lys His 450 455 460
Leu Val Ser Ala Lys Ser Asn Gly Arg Leu Thr Val Phe Pro Asp Gly
465 470 475 480
Thr Leu Glu Val Arg Tyr Ala Gln Val Gln Asp Asn Gly Thr Tyr Leu
485 490 495
Cys Ile Ala Ala Asn Ala Gly Gly Asn Asp Ser Met Pro Ala His Leu 500 505 510
His Val Arg Ser Tyr Ser Pro Asp Trp Pro His Gln Pro Asn Lys Thr 515 520 525
Phe Ala Phe Ile Ser Asn Gln Pro Gly Glu Gly Glu Ala Asn Ser Thr
530 540
Arg Ala Thr Val Pro Phe Pro Phe Asp Ile Lys Thr Leu Ile Ile Ala 545 550 555
Thr Thr Met Gly Phe Ile Ser Phe Leu Gly Val Val Leu Phe Cys Leu 565 575
Val Leu Leu Phe Leu Trp Ser Arg Gly Lys Gly Asn Thr Lys His Asn 580 585
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			LUKKI	CIEL) , 3 = (JOEIN	_E LJ	[21T]	AQ ÉL	COM .). _ _	01.2			.01 20	
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85 90 95
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Glu Arg Arg Ser His Cys Thr Leu Glu Asn Glu Pro Leu Arg Gly Phe Ser Trp Leu Ser Ser Asp Pro Gly Gly Leu Glu Ser Asp Thr Leu 125 Gln Trp Val Glu Glu Pro Gln Arg Ser Cys Thr Ala Arg Arg Cys Ala 130 135 140 Leu Gln Ala Thr Gly Gly Val Glu Pro Ala Gly Trp Lys Glu Met 150 155 160 Arg Cys His Leu Arg Ala Asn Gly Tyr Leu Cys Lys Tyr Gln Phe Glu 165 170 175 Val Leu Cys Pro Ala Pro Arg Pro Gly Ala Ala Ser Asn Leu Ser Tyr 180 185 190 Arg Ala Pro Phe Gln Leu His Ser Ala Ala Leu Asp Phe Ser Pro Pro 195 200 205 Gly Thr Glu Val Ser Ala Leu Cys Arg Gly Gln Leu Pro Ile Ser Val 210 215 220 Thr Cys Ile Ala Asp Glu Ile Gly Ala Arg Trp Asp Lys Leu Ser Gly 225 230 235 Asp Val Leu Cys Pro Cys Pro Gly Arg Tyr Leu Arg Ala Gly Lys Cys 245 250 255 Ala Glu Leu Pro Asn Cys Leu Asp Asp Leu Gly Gly Phe Ala Cys Glu 260 265 270 Cys Ala Thr Gly Phe Glu Leu Gly Lys Asp Gly Arg Ser Cys Val Thr 275 280 285 Ser Gly Glu Gly Gln Pro Thr Leu Gly Gly Thr Gly Val Pro Thr Arg 290 295 300 Arg Pro Pro Ala Thr Ala Thr Ser Pro Val Pro Gln Arg Thr Trp Pro 305 310 315 320 320 Ile Arg Val Asp Glu Lys Leu Gly Glu Thr Pro Leu Val Pro Glu Gln 325 330 335 Asp Asn Ser Val Thr Ser Ile Pro Glu Ile Pro Arg Trp Gly Ser Gln 340 345 350 Ser Thr Met Ser Thr Leu Gln Met Ser Leu Gln Ala Glu Ser Lys Ala 355 360 365 Thr Ile Thr Pro Ser Gly Ser Val Ile Ser Lys Phe Asn Ser Thr Thr 370 380 Ser Ser Ala Thr Pro Gln Ala Phe Asp Ser Ser Ser Ala Val Phe 385 390 395 400 Ile Phe Val Ser Thr Ala Val Val Leu Val Ile Leu Thr Met Thr Val Leu Gly Leu Val Lys Leu Cys Phe His Glu Ser Pro Ser Ser Gln
420 430

Pro Ar	g Lys 435	CORRI Glu	Ser	SEC Met	QUENC Gly	CE LI Pro 440	ESTIN Pro	NG FF Gly	ROM : Leu	10466 Glu	5.14 Ser 445	01.2 Asp	29.02 Pro	2P1618 Glu	3P2C1.txt
Pro Al 45		Leu	Glу	Ser	Ser 455	Ser	Ala	ніѕ	Cys	Thr 460	Asn	Asn	Gly	^val	
Lys Va 465	l Gly	Asp	Cys	Asp 470	Leu	Arg	Asp	Arg	Ala 475	Glu	Gly	Ala	Leu	Leu 480	
Ala Gl	u Ser	Pro	Leu 485	Gly	Ser	Ser	Asp	Ala 490							
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<220> <223>	Descr oligo	ipti nucl	on o	f Ari de pi	tifi robe	cial	Séq	uence	e: S	ynth	etic				
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<220> <223>	Descr oligo						Seq	uenc	e: S	ynth	etic				
<400> tgacca		ggaa	ggac	ag											20
<210> <211> <212> <213>	20 DNA	icia	.1 Se	quen	ce						,				
<220> <223>	Descr oligo	ipti nucl	on o eoti	f Ar de p	tifi robe	cial	Seq	uenc	e: S	ynth	etic				
<400> acagag		gggt	gcct	tg											20
<210> <211> <212> <213>	24 DNA	icia	₁1 Se	quen	ce										+ 2
<220> <223>	Descr oligo	ipti nucl	on o eoti	f Ar de p	tifi robe	cial	Seq	uenc	e: S	ynth	etic				·
<400> tcaggg		gtgg	tgtc	tc t	ccc										24
<210><211><211>	24														

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
          oligonucleotide probe
<400> 101
                                                                                                              24
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<210> 102
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
          oligonucleotide probe
<400> 102
                                                                                                               50 -
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<210> 103
<211> 2026
<212> DNA
<213> Homo sapiens
<400> 103
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agcgtcgagt cagacggcac cataatcgcc tttaaaagtg cctccgccct gccggccgcg 120 tatccccgg ctacctgggc cgccccgcgg cggtgcgcgc gtgagaggga gcgcgcgggc 180
agccgagcgc cggtgtgagc cagcgctgct gccagtgtga gcggcggtgt gagcgcggtg 240 ggtgcggagg ggcgtgtgtg ccggcgcgcg cgccgtgggg tgcaaacccc gagcgtctac 300 gctgccatga ggggcgcgaa cgcctgggcg ccactctgcc tgctgctggc tgccgccacc 360
cagctctcgc ggcagcagtc cccagagaga cctgttttca catgtggtgg cattcttact 420 ggagagtctg gatttattgg cagtgaaggt tttcctggag tgtaccctcc aaatagcaaa 480 tgtacttgga aaatcacagt tcccgaagga aaagtagtcg ttctcaattt ccgattcata 540
gacctcgaga gtgacaacct gtgccgctat gactttgtgg atgtgtacaa tggccatgcc 600
aatggccagc gcattggccg ctictgtggc actttccggc ctggagccct tgtgtccagt 660
ggcaacaaga tgatggtgca gatgatttct gatgccaaca cagctggcaa tggcttcatg 720 gccatgttct ccgctgctga accaaacgaa agaggggatc agtattgtgg aggactcctt 780
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ggggaagtca acgatgctag aagaattgga aagtattgtg gtgatagtcc acctgcgcca 1020 attgtgtctg agagaaatga acttcttatt cagttttat cagacttaag tttaactgca 1080 gatgggttta ttggtcacta catattcagg ccaaaaaaac tgcctacaac tacagaacag 1140
cctgicacca ccacattccc tgtaaccacg ggtttaaaac ccaccgtggc cttgigtcaa 1200
caaaagtgta gacggacggg gactctggag ggcaattatt gttcaagtga ctttgtatta 1260 gccggcactg ttatcacaac catcactcgc gatgggagtt tgcacgccac agtctcgatc 1320
atcaacatct acaaagaggg aaatttggcg attcagcagg cgggcaagaa catgagtgcc 1380 aggctgactg tcgtctgcaa gcagtgccct ctcctcagaa gaggtctaaa ttacattatt 1440
atgggccaag taggtgaaga tgggcgaggc aaaatcatgc caaacagctt tatcatgatg 1500
ttcaagacca agaatcagaa gctcctggat gccttaaaaa ataagcaatg ttaacagtga 1560 actgtgtcca tttaagctgt attctgccat tgcctttgaa agatctatgt tctctcagta 1620 gaaaaaaaaa tacttataaa attacatatt ctgaaagagg attccgaaag atgggactgg 1680
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tgctgtcaga ggagcagcta tctgattgga aacctgccga cttagtgcgg tgataggaag 1800 ctaaaagtgt caagcgttga cagcttggaa gcgtttattt atacatctct gtaaaaggat 1860 attttagaat tgagttgtgt gaagatgtca aaaaaagatt ttagaagtgc aatatttata 1920
gtgttatttg titcacctic aagcctitgc cctgaggtgt tacaatctig tcttgcgttt 1980
tctaaatcaa tgcttaataa aatatttta aaggaaaaaa aaaaaa
```

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

<211> 415 <212> PRT

<213> Homo sapiens

<400> 104 Met Arg Gly Ala Asn Ala Trp Ala Pro Leu Cys Leu Leu Leu Ala Ala 10 15 Ala Thr Gln Leu Ser Arg Gln Gln Ser Pro Glu Arg Pro Val Phe Thr 20 25 30 Cys Gly Gly Ile Leu Thr Gly Glu Ser Gly Phe Ile Gly Ser Glu Gly 35 40 Phe Pro Gly Val Tyr Pro Pro Asn Ser Lys Cys Thr Trp Lys Ile Thr 50 60 Val Pro Glu Gly Lys Val Val Val Leu Asn Phe Arg Phe Ile Asp Leu 65 70 75 80 Glu Ser Asp Asn Leu Cys Arg Tyr Asp Phe Val Asp Val Tyr Asn Gly
85 90 95 His Ala Asn Gly Gln Arg Ile Gly Arg Phe Cys Gly Thr Phe Arg Pro 100 105 110 Gly Ala Leu Val Ser Ser Gly Asn Lys Met Met Val Gln Met Ile Ser 115 120 125 Asp Ala Asn Thr Ala Gly Asn Gly Phe Met Ala Met Phe Ser Ala Ala 130 140 Glu Pro Asn Glu Arg Gly Asp Gln Tyr Cys Gly Gly Leu Leu Asp Arg 145 150 155 160 Pro Ser Gly Ser Phe Lys Thr Pro Asn Trp Pro Asp Arg Asp Tyr Pro 165 170 175 Ala Gly Val Thr Cys Val Trp His Ile Val Ala Pro Lys Asn Gln Leu 180 185 190 Ile Glu Leu Lys Phe Glu Lys Phe Asp Val Glu Arg Asp Asn Tyr Cys 195 200 205 Arg Tyr Asp Tyr Val Ala Val Phe Asn Gly Gly Glu Val Asn Asp Ala 210 215 220 Arg Arg Ile Gly Lys Tyr Cys Gly Asp Ser Pro Pro Ala Pro Ile Val 225 230 235 240 Ser Glu Arg Asn Glu Leu Leu Ile Gln Phe Leu Ser Ásp Leu Ser Leu 245 250 255 Thr Ala Asp Gly Phe Ile Gly His Tyr Ile Phe Arg Pro Lys Lys Leu 260 265 270 Pro Thr Thr Glu Gln Pro Val Thr Thr Phe Pro Val Thr Thr 275 280 285 Gly Leu Lys Pro Thr Val Ala Leu Cys Gln Gln Lys Cys Arg Arg Thr 290 295 300 Gly Thr Leu Glu Gly Asn Tyr Cys Ser Ser Asp Phe Val Leu Ala Gly

```
CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
                                         315
305
                    310
Thr Val Ile Thr Thr Ile Thr Arg Asp Gly Ser Leu His Ala Thr Val
Ser Ile Ile Asn Ile Tyr Lys Glu Gly Asn Leu Ala Ile Gln Gln Ala
340 345 350
Gly Lys Asn Met Ser Ala Arg Leu Thr Val Val Cys Lys Gln Cys Pro
Leu Leu Arg Arg Gly Leu Asn Tyr Ile Ile Met Gly Gln Val Gly Glu
Asp Gly Arg Gly Lys Ile Met Pro Asn Ser Phe Ile Met Met Phe Lys
385
Thr Lys Asn Gln Lys Leu Leu Asp Ala Leu Lys Asn Lys Gln Cys
                405
<210> 105
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 105
                                                                     22
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<210> 106
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 106
                                                                     22
gtcaaggagt cctccacaat ac
<210> 107
<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 107
                                                                     45
gtgtacaatg gccatgccaa tggccagcgc attggccgct tctgt
<210> 108
<211> 1838
<212> DNA
<213> Homo sapiens
<400> 108
cggacgcgtg ggcggacgcg tgggcggccc acggcgcccg cgggctgggg cggtcgcttc 60
                                        Page 54
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
ttccttctcc gtggcctacg agggtcccca gcctgggtaa agatggcccc atggcccccg 120
aagggcctag tcccagctgt gctctggggc ctcagcctct tcctcaacct cccaggacct
atčtěgeteč agecetetěc acetečetág tettětece egecteagee ceatčegtgt 240
catacctgcc ggggactggt tgacagcttt aacaagggcc tggagagaac catccgggac
                                                                                                    300
aactttggag gtggaaacac tgcctgggag gaagagaatt tgtccaaata caaagacagt gagacccgcc tggtagaggt gctggagggt gtgtgcagca agtcagactt cgagtgccac
                                                                                                    360
                                                                                                    420
cgcctgctgg agctgagtga ggagctggtg gagagctggt ggtttcacaa gcagcaggag 480 gccccggacc tcttccagtg gctgtgctca gattccctga agctcgctg ccccgcaggc 540 accttcgggc cctcctgct tccctgtcct ggggggaacag agaggccctg cggtggctac 600
gggcagtgtg aaggagaagg gacacgaggg ggcagcgggc actgtgactg ccaagccggc 660 tacgggggtg aggcctgtgg ccagtgtggc cttggctact ttgaggcaga acgcaacgcc 720
agccatctgg tatgttcggc ttgttttggc ccctgtgccc gatgctcagg acctgaggaa 780
tcaaactgtt tgcaatgcaa gaagggctgg gccctgcatc acctcaagtg tgtagacatt 840
gatgagtgtg gcacagaggg agccaactgt ggagctgacc aattctgcgt gaacactgag 900
ggctcctatg agtgccgaga ctgtgccaag gcctgcctag gctgcatggg ggcagggcca 960 ggtcgctgta agaagtgtag ccctggctat cagcaggtgg gctccaagtg tctcgatgtg 1020 gatgagtgtg agacagaggt gtgtccggga gagaacaagc agtgtgaaaa caccgagggc 1080 ggttatcgct gcatctgtgc cgagggctac aagcagatgg aaggcatctg tgtgaaggag 1140 cagatcccag agtgagaag gttcttctca gagatcccag agtgagagag 1200
cagatcccag agtcagcagg cttcttctca gagatgacag aagacgagtt ggtggtgctg 1200 cagcagatgt tctttggcat catcatctgt gcactggcca cgctggctgc taagggcgac 1260 ttggtgttca ccgccatctt cattggggct gtggcggcca tgactggcta ctggttgtca 1320
gagcgcagtg accgtgtgct ggagggcttc atcaagggca gataatcgcg gccaccacct 1380
ğtägğacčtč ctcčcácčca čgcťgčcccc agagcťťggg čtgcccťčcť gctggacact
caggacaget tggtttattt ttgagagtgg ggtaagcacc cctacctgcc ttacagagca
gcccaggtac ccaggcccgg gcagacaagg cccctggggt aaaaagtagc cctgaaggtg gataccatga gctcttcacc tggcggggac tggcaggctt cacaatgtgt gaatttcaaa
agttttcct taatggtggc tgctagagct ttggcccctg cttaggatta ggtggtcctc acaggggtgg ggccatcaca gctcctcct gccagctgca tgctgccagt tcctgttctg tgttcaccac atccccacac cccattgcca cttatttatt catctcagga aataaagaaa
                                                                                                    1680
ggtcttggaa agttaaaaaa aaaaaaaaa aaaaaaaa
<210> 109
<211> 420
<212> PRT
<213> Homo sapiens
<400> 109
Met Ala Pro Trp Pro Pro Lys Gly Leu Val Pro Ala Val Leu Trp Gly
1 5 10 15
Leu Ser Leu Phe Leu Asn Leu Pro Gly Pro Ile Trp Leu Gln Pro Ser
20 25 30
Pro Pro Pro Gln Ser Ser Pro Pro Pro Gln Pro His Pro Cys His Thr
35 40 45
Cys Arg Gly Leu Val Asp Ser Phe Asn Lys Gly Leu Glu Arg Thr Ile 50 60
Arg Asp Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Glu Asn Leu 65 70 75 80
Ser Lys Tyr Lys Asp Ser Glu Thr Arg Leu Val Glu Val Leu Glu Gly
85 90 95
Val Cys Ser Lys Ser Asp Phe Glu Cys His Arg Leu Leu Glu Leu Ser
100 105 110
Glu Glu Leu Val Glu Ser Trp Trp Phe His Lys Gln Gln Glu Ala Pro
115 120 125
Asp Leu Phe Gln Trp Leu Cys Ser Asp Ser Leu Lys Leu Cys Cys Pro
130 135 140
                                                           Page 55
```

```
CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1,txt
Ala Gly Thr Phe Gly Pro Ser Cys Leu Pro Cys Pro Gly Gly Thr Glu
145 150 155 160
Arg Pro Cys Gly Gly Tyr Gly Gln Cys Glu Gly Glu Gly Thr Arg Gly
165 170 175
Gly Ser Gly His Cys Asp Cys Gln Ala Gly Tyr Gly Glu Ala Cys
180 185 190
Gly Gln Cys Gly Leu Gly Tyr Phe Glu Ala Glu Arg Asn Ala Ser His
195 200 205
Leu Val Cys Ser Ala Cys Phe Gly Pro Cys Ala Arg Cys Ser Gly Pro 210 220
Glu Glu Ser Asn Cys Leu Gln Cys Lys Lys Gly Trp Ala Leu His His 225 230 235 240
Leu Lys Cys Val Asp Ile Asp Glu Cys Gly Thr Glu Gly Ala Asn Cys 245 250 255
Gly Ala Asp Gln Phe Cys Val Asn Thr Glu Gly Ser Tyr Glu Cys Arg
260 265 270
Asp Cys Ala Lys Ala Cys Leu Gly Cys Met Gly Ala Gly Pro Gly Arg 275 280 285
Cys Lys Lys Cys Ser Pro Gly Tyr Gln Gln Val Gly Ser Lys Cys Leu
290 295 300
Asp Val Asp Glu Cys Glu Thr Glu Val Cys Pro Gly Glu Asn Lys Gln 305 310 315 320
Cys Glu Asn Thr Glu Gly Gly Tyr Arg Cys Ile Cys Ala Glu Gly Tyr
325 330 335
Lys Gln Met Glu Gly Ile Cys Val Lys Glu Gln Ile Pro Glu Ser Ala
340 345 350
Gly Phe Phe Ser Glu Met Thr Glu Asp Glu Leu Val Val Leu Gln Gln 355 360 365
Met Phe Phe Gly Ile Ile Ile Cys Ala Leu Ala Thr Leu Ala Ala Lys 370 380
Gly Asp Leu Val Phe Thr Ala Ile Phe Ile Gly Ala Val Ala Ala Met
385 390 395 400
Thr Gly Tyr Trp Leu Ser Glu Arg Ser Asp Arg Val Leu Glu Gly Phe 405 415
Ile Lys Gly Arg
420
<210> 110
<211> 50
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic

Page 56

oligonucleotide probe

<220>

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

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<400> 110
                                                                                     50
cctggctatc agcaggtggg ctccaagtgt ctcgatgtgg atgagtgtga
<210> 111
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 111
                                                                                     22
attctgcgtg aacactgagg gc
<210> 112
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 112
                                                                                     22
atctgcttgt agccctcggc ac
<210> 113
<211> 1616
<212> DNA
<213> Homo sapiens
<220>
<221> modified_base
<222> (1461)..(1461)
<223> a, t, c or g
<400> 113
tgagaccete etgcagcett etcaagggac ageceeacte tgeetettge teetecaggg 60
cagcaccatg cagcccctgt ggctctgctg ggcactctgg gtgttgcccc tggccagccc 120 cggggccgcc ctgaccgggg agcagctcct gggcagcctg ctgcggcagc tgcagctcaa 180 agaggtgccc accctggaca gggccgacat ggaggagctg gtcatcccca cccacgtgag 240
ggcccagtac gtggccctgc tgcagcgcag ccacggggac cgctcccgcg gaaagaggtt 300
cagccagage treegagagg tggceggeag gtteetggeg ttggaggeea geacacacet 360
gctggtgttc ggcatggagc agcggctgcc gcccaacagc gagctggtgc aggccgtgct 420
gcggctcttc caggagccgg tccccaaggc cgcgctgcac aggcacgggc ggctgtcccc 480
gcgcagcgcc cgggcccggg tgaccgtcga gtggctgcgc gtccgcgacg acggctccaa 540 ccgcacctcc ctcatcgact ccaggctggt gtccgtccac gagagcggct ggaaggcctt 600 cgacgtgacc gaggccgtga acttctggca gcagctgagc cggccccggc agccgctgct 660
gctacaggtg tcggtgcaga gggagcatct gggcccgctg gcgtccggcg cccacaagct 720 ggtccgcttt gcctcgcagg gggcgccagc cgggcttggg gagccccagc tggagctgca 780 caccctggac cttggggact atggagctca gggcgactgt gaccctgaag caccaatgac 840
cgagggcacc cgctgctgcc gccaggagat gtacattgac ctgcagggga tgaagtgggc 900
acctaattīt tģcītcītcag gaaīgāgāat ctītggccāc tggagagccc ttgctcagtt 1380
ttctctattc ttattattca ctgcactata ttctaagcac ttacatgtgg agatactgta 1440
                                                 Page 57
```

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt acctgagggc agaaagccca ntgtgtcatt gtttacttgt cctgtcactg gatctgggct 1500 aaagtcctcc accaccactc tggacctaag acctggggtt aagtgtgggt tgtgcatccc caatccagat aataaagact ttgtaaaaca tgaataaaac acattttatt ctaaaa <210> 114 <211> 366 <212> PRT <213> Homo sapiens <400> 114 Met Gln Pro Leu Trp Leu Cys Trp Ala Leu Trp Val Leu Pro Leu Ala Ser Pro Gly Ala Ala Leu Thr Gly Glu Gln Leu Leu Gly Ser Leu Leu 20 25 30 Arg Gln Leu Gln Leu Lys Glu Val Pro Thr Leu Asp Arg Ala Asp Met 35 40 45 Glu Glu Leu Val Ile Pro Thr His Val Arg Ala Gln Tyr Val Ala Leu 50 60 Leu Gln Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln 65 70 75 80 Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Leu Glu Ala Ser Thr 85 90 95 His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu 100 105 110Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val Pro Lys Ala 115 120 125 Ala Leu His Arg His Gly Arg Leu Ser Pro Arg Ser Ala Arg Ala Arg 130 135 140 Val Thr Val Glu Trp Leu Arg Val Arg Asp Asp Gly Ser Asn Arg Thr 145 150 155 160 Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly Trp Lys 165 170 175 Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu Ser Arg 180 185 190 Pro Arg Gln Pro Leu Leu Gln Val Ser Val Gln Arg Glu His Leu 195 200 205 Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala Ser Gln 210 215 220 Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His Thr Leu 225 230 235 240 Asp Leu Gly Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu Ala Pro 245 250 255 Met Thr Glu Gly Thr Arg Cys Cys Arg Gln Glu Met Tyr Ile Asp Leu 260 265 270

Gln Gly Met Lys Trp Ala Glu Asn Trp Val Leu Glu Pro Pro Gly Phe 275 280 285

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
Leu Ala Tyr Glu Cys Val Gly Thr Cys Arg Gln Pro Pro Glu Ala Leu
                         295
Ala Phe Lys Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala Ser Glu
305
Thr Asp Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly Arg Thr
                                     330
Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg Val Gln Lys Cys Ser
            340
Cys Ala Ser Asp Gly Ala Leu Val Pro Arg Arg Leu Gln Pro
<210> 115
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 115
                                                                    21
aggactgcca taacttgcct g
<210> 116
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 116
                                                                    22
ataggagttg aagcagcgct gc
<210> 117
<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 117
                                                                    45
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<210> 118
<211> 1857
<212> DNA
<213> Homo sapiens
<400> 118
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aaggcgcaag tcgagaggaa actgttgtgc ctcttcatat tggcgatcct gttgtgctcc 120
ctggcattgg gcagtgttac agtgcactct tctgaacctg aagtcagaat tcctgagaat 180
aatcctgtga agttgtcctg tgcctactcg ggcttttctt ctccccgtgt ggagtggaag 240
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1,txt
tttgaccaag gagacaccac cagactcgtt tgctataata acaagatcac agcttcctat 300
gaggaccggg tgaccttctt gccaactggt atcaccttca agtccgtgac acgggaagac 360
actgggacat acacttgtat ggtctctgag gaaggcggca acagctatgg ggaggtcaag 420 gtcaagctca tcgtgcttgt gcctccatcc aagcctacag ttaacatccc ctcctctgcc 480
accattggga accgggcagt gctgacatgc tcagaacaag atggttcccc accttctgaa 540
tacacctggt tcaaagatgg gatagtgatg cctacgaatc ccaaaagcac ccgtgccttc 600 agcaactctt cctatgtcct gaatcccaca acaggagagc tggtctttga tcccctgtca 660 gcctctgata ctggagaata cagctgtgag gcacggaatg ggtatgggac acccatgact 720 tcaaatgctg tgcgcatgga agctgtggag cggaatgtgg gggtcatcgt ggcagccgtc 780
cttgtaaccc tgattctcct gggaatcttg gtttttggca tctggtttgc ctatagccga 840
ggccactttg acagaacaaa gaaagggact tcgagtaaga aggtgattta cagccagcct 900
agtgcccgaa gtgaaggaga attcaaacag acctcgtcat tcctggtgtg agcctggtcg 960
gčtčaccýce ťaťcaťčtýc atttycettá etcagýtyct acegýáctet gycecétyať 1020
gtctgtagtt tcacaggatg ccttatttgt cttctacacc ccacagggcc ccctacttct 1080
tcggatgtgt ttttaataat gtcagctatg tgccccatcc tccttcatgc cctccctccc 1140 tttcctacca ctgctgagtg gcctggaact tgtttaaagt gtttattccc catttctttg 1200 agggatcagg aaggaatcct gggtatgcca ttgacttccc ttctaagtag acagcaaaaa 1260 tggcgggggt cgcaggaatc tgcactcaac tgcccacctg gctggcaggg atctttgaat 1320 aggtatcttg agcttgtt tgggctcttt ccttgtgatt tgacgaccag ggccagctgt 1380
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<212> PRT
<213> Homo sapiens
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Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro Val Lys Leu
35 40 45
Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe 50 55 60
Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr 65 70 75 80
Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe 85 90 95
Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser
100 105 110
Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val
115 120 125
Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr
130 135 140
Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro
                                                             155
                                                           Page 60
```

```
CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn
                 165
Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro
Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly 195 200 205
Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser 210 220
Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val
225 230 235 240
Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly
Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys
Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala Arg Ser Glu 275 280 285
Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val
    290
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<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 120
                                                                       24
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<210> 121
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 121
                                                                       50
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<210> 122
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 122
                                                                       20
acacctggtt caaagatggg
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

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 <211> 24
 <212> DNA
 <213> Artificial Sequence
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           oligonucleotide probe
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                                                                                                                 24
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 <211> 20
 <212> DNA
 <213> Artificial Sequence
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          oligonucleotide probe
 <400> 124
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                                                                                                                 20
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<211> 1210
 <212> DNA
 <213> Homo sapiens
<400> 126
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ggcgctgcga cggccaccca gactgtcccg actccagcga cgagctcggc tgtggaacca 540
atgagateet eeeggaaggg gatgeeacaa eeatggggee eeetgtgaee etggagagtg 600
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agctaggatg gggaacctgc cacagccaga actgaggggc tggccccagg cagctcccag 1140
ggggtágaac ggccctgtgc ttaagacact ccctgctgcc ccgtctgagg gtggccgatta 1200
                                                                 Page 62
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt aagttgcttc

<210> 127 <211> 282

<212> PRT

<213> Homo sapiens

<400> 127

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Ala Ala Ser Pro Leu Ser Thr Pro Thr Ser Ala Gln Ala Ala Gly
35 40 45

Pro Ser Ser Gly Ser Cys Pro Pro Thr Lys Phe Gln Cys Arg Thr Ser 50 55 60

Gly Leu Cys Val Pro Leu Thr Trp Arg Cys Asp Arg Asp Leu Asp Cys 65 70 75 80

Ser Asp Gly Ser Asp Glu Glu Glu Cys Arg Ile Glu Pro Cys Thr Gln
85 90 95

Lys Gly Gln Cys Pro Pro Pro Pro Gly Leu Pro Cys Pro Cys Thr Gly 100 105 110

Val Ser Asp Cys Ser Gly Gly Thr Asp Lys Lys Leu Arg Asn Cys Ser 115 120 125

Arg Leu Ala Cys Leu Ala Gly Glu Leu Arg Cys Thr Leu Ser Asp Asp 130 140

Cys Ile Pro Leu Thr Trp Arg Cys Asp Gly His Pro Asp Cys Pro Asp 145 150 155 160

Ser Ser Asp Glu Leu Gly Cys Gly Thr Asn Glu Ile Leu Pro Glu Gly 165 170 175

Asp Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val Thr Ser 180 185 190

Leu Arg Asn Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val 195 200 205

Pro Ser Val Gly Asn Ala Thr Ser Ser Ser Ala Gly Asp Gln Ser Gly 210 215 220

Ser Pro Thr Ala Tyr Gly Val Ile Ala Ala Ala Ala Val Leu Ser Ala 225 230 235 240

Ser Leu Val Thr Ala Thr Leu Leu Leu Leu Ser Trp Leu Arg Ala Gln 245 250 255

Glu Arg Leu Arg Pro Leu Gly Leu Leu Val Ala Met Lys Glu Ser Leu 260 265 270

Leu Leu Ser Glu Gln Lys Thr Ser Leu Pro 275 280

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
          oligonucleotide probe
<400> 128
                                                                                                                24
aagttccagt gccgcaccag tggc
<210> 129
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
          oligonucleotide probe
<400> 129
ttggttccac agccgagctc gtcg
                                                                                                                24
<210> 130
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
          oligonucleotide probe
<400> 130
                                                                                                                50
qaqqaqqaqt gcaggattga gccatgtacc cagaaagggc aatgcccacc
<210> 131
<211> 1843
<212> DNA
<213> Homo sapiens
<220>
<221> modified_base
<222> (1837)..(1837)
<223> a, t, c or g
<400> 131
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cagactettg caagetggat geetetgtg gatgaaagat gtateatgga atgaaceega 180 geaatggaga tggatteta gageageage ageageagea geaaceeteag tececeaga 240 gaetettgge egtgateetg tggtteage tggegetgtg etteggeet geaaggetea 300 egggegggtt egatgaeett caagtgtgtg etgaeeeegg catteeegg aatggettea 360 ggaeeeeeag eggaggggtt teettggag getetgage eegatteae tgeaagaegg 420 gatteaaget gaagggeget acaaagagae tgtgtttgaa geattttaat ggaaceeetag 480 geteggateee aagtgataat teestetgg tgeaagaaga ttgeaagaaga tgeaagaaga eggaaceeetag 540
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atgaaggatt caagatccgg taccccgacc tacacaatat ggtttcatta tgtcgcgatg 660
atggaacgtg gaataatctg cccatctgtc aaggctgcct gagacctcta gcctcttcta 720 atggctatgt aaacatctct gagctccaga cctccttccc ggtggggact gtgatctcct 780 atcgctgctt tcccggattt aaacttgatg ggtctgcgta tcttgagtgc ttacaaaaacc 840 ttatctggtc gtccagccca ccccggtgcc ttgctctgga agcccaagtc tgtccactac 900 ctccaatggt gagtcacgga gatttcgtct gccacccgcg gccttgtgag cgctacaacc 960 acggaactgt ggtggagttt tactgcgatc ctggctacaag cctccaccagc gactacaagt 1020
                                                                 Page 64
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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cğgcaaccağ tgtgctgctg gtgctgctgc tcgtcatcct ggccaggatg ttccagacca 1200
agttcaaggc ccactttccc cccagggggc ctccccggag ttccagcagt gaccctgact 1260
tīgtggtggt agacggcgtg cccgīcātgc tcccgtccta tgacgaagct gtgagtggcg
gcitgagige citaggecee gggtacatgg ectetgtggg ceagggetge ecettaeceg
tggacgacca gagccccca gcataccccg gctcagggga cacggacaca ggcccagggg
agtcagaaac ctgtgacagc gtctcaggct cttctgagct gctccaaagt ctgtattcac
                                                                            1500
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cggcagagga ggtggcatcc accagcccag gcatccatca tgcccactgg gtgttgttcc taagaaactg attgattaaa aaatttccca aagtgtcctg aagtgtctct tcaaatacat
gttgatctgt ggagttgatt cctttccttc tcttggtttt agacaaatgt aaacaaagct 1740
čtgatcctia aaattgctat gctgatagag tggtgagggc tggaagctig atcaagtcct 1800
gtītcttctt gacacāgact gatītaaaāaī tāāaāgnāāa aāā
                                                                             1843
<210> 132
<211> 490
<212> PRT
<213> Homo sapiens
<400> 132
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Gln Gln Gln Gln Gln Pro Gln Ser Pro Gln Arg Leu Leu Ala Val
20 25 30
Ile Leu Trp Phe Gln Leu Ala Leu Cys Phe Gly Pro Ala Gln Leu Thr 35 40 45
Gly Gly Phe Asp Asp Leu Gln Val Cys Ala Asp Pro Gly Ile Pro Glu 50 60
Asn Gly Phe Arg Thr Pro Ser Gly Gly Val Phe Phe Glu Gly Ser Val
65 70 75 80
Ala Arg Phe His Cys Gln Asp Gly Phe Lys Leu Lys Gly Ala Thr Lys
85 90 95
Arg Leu Cys Leu Lys His Phe Asn Gly Thr Leu Gly Trp Ile Pro Ser
100 105 110
Asp Asn Ser Ile Cys Val Gln Glu Asp Cys Arg Ile Pro Gln Ile Glu
115 120 125
Asp Ala Glu Ile His Asn Lys Thr Tyr Arg His Gly Glu Lys Leu Ile
130 135 140
Ile Thr Cys His Glu Gly Phe Lys Ile Arg Tyr Pro Asp Leu His Asn
145 150 155 160
Met Val Ser Leu Cys Arg Asp Asp Gly Thr Trp Asn Asn Leu Pro Ile
165 170 175
Cys Gln Gly Cys Leu Arg Pro Leu Ala Ser Ser Asn Gly Tyr Val Asn
180 185 190
Ile Ser Glu Leu Gln Thr Ser Phe Pro Val Gly Thr Val Ile Ser Tyr
195 200 205
Arg Cys Phe Pro Gly Phe Lys Leu Asp Gly Ser Ala Tyr Leu Glu Cys 210 220
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
Leu Gln Asn Leu Ile Trp Ser Ser Ser Pro Pro Arg Cys Leu Ala Leu 225 230 235 240
Glu Ala Gln Val Cys Pro Leu Pro Pro Met Val Ser His Gly Asp Phe 245 250 255
Val Cys His Pro Arg Pro Cys Glu Arg Tyr Asn His Gly Thr Val Val 260 265 270
Glu Phe Tyr Cys Asp Pro Gly Tyr Ser Leu Thr Ser Asp Tyr Lys Tyr 275 280 285
Ile Thr Cys Gln Tyr Gly Glu Trp Phe Pro Ser Tyr Gln Val Tyr Cys 290 295 300
Ile Lys Ser Glu Gln Thr Trp Pro Ser Thr His Glu Thr Leu Leu Thr 305 310 315
Thr Trp Lys Ile Val Ala Phe Thr Ala Thr Ser Val Leu Leu Val Leu
Leu Leu Val Ile Leu Ala Arg Met Phe Gln Thr Lys Phe Lys Ala His 340 345 350
Phe Pro Pro Arg Gly Pro Pro Arg Ser Ser Ser Ser Asp Pro Asp Phe 355 360 365
Val Val Asp Gly Val Pro Val Met Leu Pro Ser Tyr Asp Glu Ala
370 375 380
Val Ser Gly Gly Leu Ser Ala Leu Gly Pro Gly Tyr Met Ala Ser Val
385 390 395 400
Gly Gln Gly Cys Pro Leu Pro Val Asp Asp Gln Ser Pro Pro Ala Tyr
405 410 415
Pro Gly Ser Gly Asp Thr Asp Thr Gly Pro Gly Glu Ser Glu Thr Cys 420 430
Asp Ser Val Ser Gly Ser Ser Glu Leu Leu Gln Ser Leu Tyr Ser Pro
Pro Arg Cys Gln Glu Ser Thr His Pro Ala Ser Asp Asn Pro Asp Ile
450 460
Ile Ala Ser Thr Ala Glu Glu Val Ala Ser Thr Ser Pro Gly Ile His
465 470 475 480
His Ala His Trp Val Leu Phe Leu Arg Asn
<210> 133
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 133
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atctcctatc gctgctttcc cgg

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<210> 134
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
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           oligonucleotide probe
<400> 134
                                                                                                                              23
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<211> 50
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
           oligonucleotide probe
<400> 135
                                                                                                                               50
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<211> 1815
<212> DNA
<213> Homo sapiens
<400> 136
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ccgtagcgcc cgagtgtcgg ggggcgcacc cgagtcgggc catgaggccg ggaaccgcgc 180 tacaggccgt gctgctggcc gtgctgctgg tggggctgcg ggccgcgacg ggtcgcctgc 240 tgagtgcctc ggatttggac ctcagaggag ggcagccagt ctgccgggga gggacacaga 300 ggccttgtta taaagtcatt tacttccatg atacttctcg aagactgaac tttgaggaag 360 ccaaagaagc ctgcaggagg gatggaggcc agctagtcag catcgagtct gaagatgaac 420 agaaactgat agaaaagttc attgaaaacc tcttgccatc tgatggtgac ttctggattg 480 ggctcaggag gcgtgaggag aaacaaagca atagcacagc ctgccaggac ctttatgctt 540 ggactgatgg cagcatatca caatttagga actggtatgt ggatgagccg tcctgcggca 600 gcgaggtctg cgtgatga taccatcagc catcgagtcc cgctgccatc ggaggcccc 660
gcgaggtctg cgtggtcatg taccatcagc catcggcacc cgctggcatc ggaggcccct 660
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cacctgtact tccagaagaa acacaggaag aagatgccaa aaaaacattt aaagaaagta 840
gagaagctgc cttgaatctg gcctacatcc taatccccag cattcccctt ctcctccc 900 ttgtggtcac cacagttgta tgttgggttt ggatctgtag aaaaagaaaa cgggagcagc 960 cagacctag cacaaagaag caacacacca tctggccctc tcctcaccag ggaaacagcc 1020 cggacctaga ggtctacaat gtcataagaa aacaaagcga agctgactta gctgagaccc 1080 ggccagacct gaagaatatt tcattccgag tgtgttcggg agaagccact cccgatgaca 1140
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ttggcctgtg catcggcaat tctcatatct gtttttttca aagaataaaa tcaaataaag 1800
agcaggaaaa aaaaa
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<211> 382 <212> PRT

<213> Homo sapiens

<400> 137 Met Arg Pro Gly Thr Ala Leu Gln Ala Val Leu Leu Ala Val Leu Leu 1 10 15 Val Gly Leu Arg Ala Ala Thr Gly Arg Leu Leu Ser Ala Ser Asp Leu
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165 170 175 Asn Met Lys Asn Asn Phe Ile Cys Lys Tyr Ser Asp Glu Lys Pro Ala 180 185 190 val Pro Ser Arg Glu Ala Glu Gly Glu Glu Thr Glu Leu Thr Thr Pro 195 200 205 Val Leu Pro Glu Glu Thr Gln Glu Glu Asp Ala Lys Lys Thr Phe Lys 210 220 Glu Ser Arg Glu Ala Ala Leu Asn Leu Ala Tyr Ile Leu Ile Pro Ser 225 230 235 240 Ile Pro Leu Leu Leu Leu Val Val Thr Thr Val Val Cys Trp Val 245 250 255 Trp Ile Cys Arg Lys Arg Lys Arg Glu Gln Pro Asp Pro Ser Thr Lys 260 265 270 Lys Gln His Thr Ile Trp Pro Ser Pro His Gln Gly Asn Ser Pro Asp 285 Leu Glu Val Tyr Asn Val Ile Arg Lys Gln Ser Glu Ala Asp Leu Ala 290 295 300 Glu Thr Arg Pro Asp Leu Lys Asn Ile Ser Phe Arg Val Cys Ser Gly Page 68

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
                                                                         320
305
                        310
Glu Ala Thr Pro Asp Asp Met Ser Cys Asp Tyr Asp Asn Met Ala Val
Asn Pro Ser Glu Ser Gly Phe Val Thr Leu Val Ser Val Glu Ser Gly
Phe Val Thr Asn Asp Ile Tyr Glu Phe Ser Pro Asp Gln Met Gly Arg
                                  360
Ser Lys Glu Ser Gly Trp Val Glu Asn Glu Ile Tyr Gly Tyr
<210> 138
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 138
                                                                                 50
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<210> 139
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 139
                                                                                 24
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<210> 140
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 140
                                                                                 24
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<210> 141
<211> 1514
<212> DNA
<213> Homo sapiens
<400> 141
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tčeteteaag eegeteege tggaggagea ggtagagtgg aacceecage tattagaggt 420
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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cctcccaagc tgccccatgc tgaagagatt tgcacggatg atcgaacaga gagctgtgga 1020
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                                                                            1080
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35 40 45
Lys Leu Lys Met Val Gln Val Val Phe Arg His Gly Ala Arg Ser Pro 50 60
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Leu Glu Val Pro Pro Gln Thr Gln Phe Asp Tyr Thr Val Thr Asn Leu
85 90 95
Ala Gly Gly Pro Lys Pro Tyr Ser Pro Tyr Asp Ser Gln Tyr His Glu
100 105 110
Thr Thr Leu Lys Gly Gly Met Phe Ala Gly Gln Leu Thr Lys Val Gly 115 120 125
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130 135 140
Glu Asp Ile Pro Phe Leu Ser Pro Thr Phe Asn Pro Gln Glu Val Phe
145 150 155 160
Ile Arg Ser Thr Asn Ile Phe Arg Asn Leu Glu Ser Thr Arg Cys Leu
165 170 175
Leu Ala Gly Leu Phe Gln Cys Gln Lys Glu Gly Pro Ile Ile Ile His
180 185 190
Thr Asp Glu Ala Asp Ser Glu Val Leu Tyr Pro Asn Tyr Gln Ser Cys
                                             Page 70
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
                                200
Trp Ser Leu Arg Gln Arg Thr Arg Gly Arg Arg Gln Thr Ala Ser Leu 210 220
Gln Pro Gly Ile Ser Glu Asp Leu Lys Lys Val Lys Asp Arg Met Gly 225 230 235
Ile Asp Ser Ser Asp Lys Val Asp Phe Phe Ile Leu Leu Asp Asn Val
245 250 255
Ala Ala Glu Gln Ala His Asn Leu Pro Ser Cys Pro Met Leu Lys Arg
260 265 270
Phe Ala Arg Met Ile Glu Gln Arg Ala Val Asp Thr Ser Leu Tyr Ile
275 280 285
Leu Pro Lys Glu Asp Arg Glu Ser Leu Gln Met Ala Val Gly Pro Phe
290 295 300
Leu His Ile Leu Glu Ser Asn Leu Leu Lys Ala Met Asp Ser Ala Thr
305 310 315 320
305
Ala Pro Asp Lys Ile Arg Lys Leu Tyr Leu Tyr Ala Ala His Asp Val
325 330 335
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Pro Pro Phe Ala Val Asp Leu Thr Met Glu Leu Tyr Gln His Leu Glu
Ser Lys Glu Trp Phe Val Gln Leu Tyr Tyr His Gly Lys Glu Gln Val
370 375 380
Pro Arg Gly Cys Pro Asp Gly Leu Cys Pro Leu Asp Met Phe Leu Asn 385 390 395
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420 425
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CORRECTED SEQUENCE LISTING FROM $10466.14\ 01.29.02P1618P2C1.txt$ oligonucleotide probe

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt

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<213> Homo sapiens

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35 40 45 Cys Asp Asp Gly Trp Asp Ile Lys Asp Val Ala Val Leu Cys Arg Glu 50 60 Leu Gly Cys Gly Ala Ala Ser Gly Thr Pro Ser Gly Ile Leu Tyr Glu 65 70 75 80 Pro Pro Ala Glu Lys Glu Gln Lys Val Leu Ile Gln Ser Val Ser Cys 85 90 95 Thr Gly Thr Glu Asp Thr Leu Ala Gln Cys Glu Gln Glu Glu Val Tyr 100 105 110 Asp Cys Ser His Asp Glu Asp Ala Gly Ala Ser Cys Glu Asn Pro Glu 115 120 125 Ser Ser Phe Ser Pro Val Pro Glu Gly Val Arg Leu Ala Asp Gly Pro 130 140 Gly His Cys Lys Gly Arg Val Glu Val Lys His Gln Asn Gln Trp Tyr 145 150 155 160 Thr Val Cys Gln Thr Gly Trp Ser Leu Arg Ala Ala Lys Val Val Cys 165 170 175 Arg Gln Leu Gly Cys Gly Arg Ala Val Leu Thr Gln Lys Arg Cys Asn 180 185 190 Lys His Ala Tyr Gly Arg Lys Pro Ile Trp Leu Ser Gln Met Ser Cys 195 200 205 Ser Gly Arg Glu Ala Thr Leu Gln Asp Cys Pro Ser Gly Pro Trp Gly 210 215 220 Lys Asn Thr Cys Asn His Asp Glu Asp Thr Trp Val Glu Cys Glu Asp 225 230 235 240 Pro Phe Asp Leu Arg Leu Val Gly Gly Asp Asn Leu Cys Ser Gly Arg 245 250 255 Leu Glu Val Leu His Lys Gly Val Trp Gly Ser Val Cys Asp Asp Asn 260 265 Trp Gly Glu Lys Glu Asp Gln Val Val Cys Lys Gln Leu Gly Cys Gly 275 280 285 Lys Ser Leu Ser Pro Ser Phe Arg Asp Arg Lys Cys Tyr Gly Pro Gly

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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gctcatcaga gaacttaccg cttctcatgc caccaaggtg cagacacaca agccttactt 420
ggtgaccttc gacctcacag actctggggc catagttgca gcagcagctg agatcctgca 480
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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ğgtgāccgtc atcagccccg gctacatcca caccaacctc tctgtaaatg ccatcaccgc 840
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Ala Tyr Leu Arg Asn Ala Val Val Val Ile Thr Gly Ala Thr Ser Gly
35 40 45
Leu Gly Lys Glu Cys Ala Lys Val Phe Tyr Ala Ala Gly Ala Lys Leu 50 60
Val Leu Cys Gly Arg Asn Gly Gly Ala Leu Glu Glu Leu Ile Arg Glu
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Leu Thr Ala Ser His Ala Thr Lys Val Gln Thr His Lys Pro Tyr Leu
85 90 95
Val Thr Phe Asp Leu Thr Asp Ser Gly Ala Ile Val Ala Ala Ala Ala 100 105 110
Glu Ile Leu Gln Cys Phe Gly Tyr Val Asp Ile Leu Val Asn Asn Ala
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Gly Ile Ser Tyr Arg Gly Thr Ile Met Asp Thr Thr Val Asp Val Asp 130 140
Lys Arg Val Met Glu Thr Asn Tyr Phe Gly Pro Val Ala Leu Thr Lys
145 150 155 160
Ala Leu Leu Pro Ser Met Ile Lys Arg Arg Gln Gly His Ile Val Ala
165 170 175
Ile Ser Ser Ile Gln Gly Lys Met Ser Ile Pro Phe Arg Ser Ala Tyr
180 185 190
Ala Ala Ser Lys His Ala Thr Gln Ala Phe Phe Asp Cys Leu Arg Ala
195 200 205
Glu Met Glu Gln Tyr Glu Ile Glu Val Thr Val Ile Ser Pro Gly Tyr
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and the control of th	•
CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P Ile His Thr Asn Leu Ser Val Asn Ala Ile Thr Ala Asp Gly Ser Arg 225 230 235 240	2C1.txt
Tyr Gly Val Met Asp Thr Thr Ala Gln Gly Arg Ser Pro Val Glu 245 250 255	•
Val Ala Gln Asp Val Leu Ala Ala Val Gly Lys Lys Lys Asp Val 260 265 270	
Ile Leu Ala Asp Leu Leu Pro Ser Leu Ala Val Tyr Leu Arg Thr Leu 275 280 285	
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Ser Val Thr Gly Glu Ile Val Leu Ile Thr Gly Ala Gly His Gly Ile
35 40 45
Gly Arg Leu Thr Ala Tyr Glu Phe Ala Lys Leu Lys Ser Lys Leu Val
Leu Trp Asp Ile Asn Lys His Gly Leu Glu Glu Thr Ala Ala Lys Cys
65 70 75 80
Lys Gly Leu Gly Ala Lys Val His Thr Phe Val Val Asp Cys Ser Asn 85 90 95
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Asp Val Ser Ile Leu Val Asn Asn Ala Gly Val Val Tyr Thr Ser Asp
115 120 125
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130 135 140
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150 155 160
Lys Asn Asn His Gly His Ile Val Thr Val Ala Ser Ala Ala Gly His
165 170 175
Val Ser Val Pro Phe Leu Leu Ala Tyr Cys Ser Ser Lys Phe Ala Ala
180 185 190
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195 200 205
Thr Gly Val Lys Thr Thr Cys Leu Cys Pro Asn Phe Val Asn Thr Gly 210 220
Phe Ile Lys Asn Pro Ser Thr Ser Leu Gly Pro Thr Leu Glu Pro Glu 225 230
Glu Val Val Asn Arg Leu Met His Gly Ile Leu Thr Glu Gln Lys Met
245 250 255
Ile Phe Ile Pro Ser Ser Ile Ala Phe Leu Thr Thr Leu Glu Arg Ile
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Pro Gly Pro Cys Asp Gly Leu Phe Arg Ser Leu Tyr Arg Ser Val Ser 20 25 30 Met Pro Pro Lys Gly Asp Ser Gly Gln Pro Leu Phe Leu Thr Pro Tyr 35 40 45 Ile Glu Ala Gly Lys Ile Gln Lys Gly Arg Glu Leu Ser Leu Val Gly 50 60 Pro Phe Pro Gly Leu Asn Met Lys Ser Tyr Ala Gly Phe Leu Thr Val 65 70 75 80 Asn Lys Thr Tyr Asn Ser Asn Leu Phe Phe Trp Phe Phe Pro Ala Gln 85 90 95 Ile Gln Pro Glu Asp Ala Pro Val Val Leu Trp Leu Gln Gly Gly Pro 100 105 110 Gly Gly Ser Ser Met Phe Gly Leu Phe Val Glu His Gly Pro Tyr Val 115 120 125 val Thr Ser Asn Met Thr Leu Arg Asp Arg Asp Phe Pro Trp Thr Thr 130 135 140 Thr Leu Ser Met Leu Tyr Ile Asp Asn Pro Val Gly Thr Gly Phe Ser 145 150 155 160 Phe Thr Asp Asp Thr His Gly Tyr Ala Val Asn Glu Asp Asp Val Ala 165 170 175 Arg Asp Leu Tyr Ser Ala Leu Ile Gln Phe Phe Gln Ile Phe Pro Glu 180 185 190 Tyr Lys Asn Asn Asp Phe Tyr Val Thr Gly Glu Ser Tyr Ala Gly Lys 195 200 205 Tyr Val Pro Ala Ile Ala His Leu Ile His Ser Leu Asn Pro Val Arg 210 215 220 Glu Val Lys Ile Asn Leu Asn Gly Ile Ala Ile Gly Asp Gly Tyr Ser 225 230 235 240 Asp Pro Glu Ser Ile Ile Gly Gly Tyr Ala Glu Phe Leu Tyr Gln Ile 245 250 255 Gly Leu Leu Asp Glu Lys Gln Lys Lys Tyr Phe Gln Lys Gln Cys His 260 265 270 Glu Cys Ile Glu His Ile Arg Lys Gln Asn Trp Phe Glu Ala Phe Glu 275 280 285 Ile Leu Asp Lys Leu Leu Asp Gly Asp Leu Thr Ser Asp Pro Ser Tyr 290 295 300 Phe Gln Asn Val Thr Gly Cys Ser Asn Tyr Tyr Asn Phe Leu Arg Cys 305 310 315 Thr Glu Pro Glu Asp Gln Leu Tyr Tyr Val Lys Phe Leu Ser Leu Pro 325 330 335 Glu Val Arg Gln Ala Ile His Val Gly Asn Gln Thr Phe Asn Asp Gly 340 345 Page 80

			C	ORRE	ECTE	SEC	QUENC	E LI	ISTIN	NG FF	ROM 1	L0466	5.14	01.2	29.02	P1618	P2Ç1.	t
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	Pro	Trp 370	Leu	Thr	Glu	Ile	Met 375	Asn	Asn	Tyr	Lys	va1 380	Leu	Ile	Tyr	Asn		
	Gly 385	Gln	Leu	Asp	Ile	11e 390	val	Ala	Ala	Ala	Leu 395	Thr	Glu	Arg	Ser	Leu 400		
	Met	Gly	Met	Asp	Trp 405	Lys	GŢy	ser	Gln	Glu 410	Tyr	Lys	Lys	Ala	Glu 415	Lys		-
	Lys	٧a٦	Trp	Lys 420	Ile	Phe	Lys	Ser	Asp 425	Ser	Glu	val	Ala	Gly 430	Tyr	Ile		
	Arg	Gln	Ala 435	Gly	Asp	Phe	His	Gln 440	∨al	Ile	Ile	Arg	Gly 445	Gly	Gly	His		
	Ile	Leu 450	Pro	Tyr	Asp	Gln	Pro 455	Leu	Arg	Ala	Phe	Asp 460	Met	Ile	Asn	Arg		
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)> 1(L> 5(,													

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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 <213> Homo sapiens
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Gly Cys Thr Arg Ala Gly Val Glu Tyr Arg Leu Leu Gln Ala Ala Tyr
355 360 365
Leu Ala Lys Pro Gly Asp Ser Leu Ala Gln Ala Phe Asn Ile Thr Ser 370 380
Gln Asp Asp Val Leu Phe Ala Ile Phe Ser Lys Gly Gln Lys Gln Tyr
385 390 395 400
His His Pro Pro Asp Asp Ser Ala Leu Cys Ala Phe Pro Ile Arg Ala
405 410 415
Ile Asn Leu Glm Ile Lys Glu Arg Leu Glm Ser Cys Tyr Glm Gly Glu
Gly Asn Leu Glu Leu Asn Trp Leu Leu Gly Lys Asp Val Gln Cys Thr
435 440 445
Lys Ala Pro Val Pro Ile Asp Asp Asn Phe Cys Gly Leu Asp Ile Asn 450 455 460
Gln Pro Leu Gly Gly Ser Thr Pro Val Glu Gly Leu Thr Leu Tyr Thr
465 470 475 480
Thr Ser Arg Asp Arg Mét Thr Ser Val Ala Ser Tyr Val Tyr Asn Gly
485 490 495
Tyr Ser Val Val Phe Val Gly Thr Lys Ser Gly Lys Leu Lys Lys Val 500 505 510
Arg Val Tyr Glu Phe Arg Cys Ser Asn Ala Ile His Leu Leu Ser Lys 515 520 525
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Leu Tyr Phe Leu Gly Glu Gln Arg
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Ser Thr Leu Val Pro Leu Arg Leu Arg His Arg Gln Leu Gly Leu Gln 35 40
Ala Lys Gly Trp Asn Phe Met Leu Glu Asp Ser Thr Phe Trp Ile Phe 50 60
Gly Gly Ser Ile His Tyr Phe Arg Val Pro Arg Glu Tyr Trp Arg Asp
65 70 75 80
Arg Leu Leu Lys Met Lys Ala Cys Gly Leu Asn Thr Leu Thr Thr Tyr 85 90 95
Val Pro Trp Asn Leu His Glu Pro Glu Arg Gly Lys Phe Asp Phe Ser
Gly Asn Leu Asp Leu Glu Ala Phe Val Leu Met Ala Ala Glu Ile Gly
115 120 125
Leu Trp Val Ile Leu Arg Pro Gly Pro Tyr Ile Cys Ser Glu Met Asp
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145 150 155 160
Arg Thr Thr Tyr Lys Gly Phe Thr Glu Ala Val Asp Leu Tyr Phe Asp
165 170 175
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt His Leu Met Ser Arg Val Val Pro Leu Gln Tyr Lys Arg Gly Gly Pro 180 185 Ile Ile Ala Val Gln Val Glu Asn Glu Tyr Gly Ser Tyr Asn Lys Asp 195 200 205 Pro Ala Tyr Met Pro Tyr Val Lys Lys Ala Leu Glu Asp Arg Gly Ile 210 215 220 Val Glu Leu Leu Eur Thr Ser Asp Asn Lys Asp Gly Leu Ser Lys Gly 225 230 235 Ile Val Gln Gly Val Leu Ala Thr Ile Asn Leu Gln Ser Thr His Glu 245 250 255 Leu Gln Leu Leu Thr Thr Phe Leu Phe Asn Val Gln Gly Thr Gln Pro 260 265 270 Lys Met Val Met Glu Tyr Trp Thr Gly Trp Phe Asp Ser Trp Gly Gly 285 Pro His Asn Ile Leu Asp Ser Ser Glu Val Leu Lys Thr Val Ser Ala 290 295 300 Ile Val Asp Ala Gly Ser Ser Ile Asn Leu Tyr Met Phe His Gly Gly 305 310 315 Thr Asn Phe Gly Phe Met Asn Gly Ala Met His Phe His Asp Tyr Lys 325 330 335 Ser Asp Val Thr Ser Tyr Asp Tyr Asp Ala Val Leu Thr Glu Ala Gly 340 345 350 Asp Tyr Thr Ala Lys Tyr Met Lys Leu Arg Asp Phe Phe Gly Ser Ile 355 360 365 Ser Gly Ile Pro Leu Pro Pro Pro Pro Asp Leu Pro Lys Met Pro 370 380 Tyr Glu Pro Leu Thr Pro Val Leu Tyr Leu Ser Leu Trp Asp Ala Leu 385 390 395 400 Lys Tyr Leu Gly Glu Pro Ile Lys Ser Glu Lys Pro Ile Asn Met Glu 405 410 415 Asn Leu Pro Val Asn Gly Gly Asn Gly Gln Ser Phe Gly Tyr Ile Leu 420 430 Tyr Glu Thr Ser Ile Thr Ser Ser Gly Ile Leu Ser Gly His Val His 435 440 445 Asp Arg Gly Gln Val Phe Val Asn Thr Val Ser Ile Gly Phe Leu Asp 450 460 Tyr Lys Thr Thr Lys Ile Ala Val Pro Leu Ile Gln Gly Tyr Thr Val 465 470 475 480 Leu Arg Ile Leu Val Glu Asn Arg Gly Arg Val Asn Tyr Gly Glu Asn 485 490 495 Ile Asp Asp Gln Arg Lys Gly Leu Ile Gly Asn Leu Tyr Leu Asn Asp

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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530 540
Pro Thr Leu Pro Ala Phe Phe Leu Gly Ser Leu Ser Ile Ser Ser Thr
545 550 555 560
Pro Cys Asp Thr Phe Leu Lys Leu Glu Gly Trp Glu Lys Gly Val Val 565 570 575
Phe Ile Asn Gly Gln Asn Leu Gly Arg Tyr Trp Asn Ile Gly Pro Gln 580 585
Lys Thr Leu Tyr Leu Pro Gly Pro Trp Leu Ser Ser Gly Ile Asn Gln 595 600
Val Ile Val Phe Glu Glu Thr Met Ala Gly Pro Ala Leu Gln Phe Thr
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Glu Thr Pro His Leu Gly Arg Asn Gln Tyr Ile Lys 625 630 635
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Glu Met Gly Gly Leu Pro Ser Trp Leu Leu Arg Lys Pro Glu Ile His
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Lys Val Leu Leu Pro Lys Ile Tyr Pro Trp Leu Tyr His Asn Gly Gly
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565 570 575 Trp Thr Lys Gln Gly Pro Gln Gln Thr Leu Tyr Val Pro Arg Phe Leu
580 585 590 Page 90

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Leu Phe Pro Arg Gly Ala Leu Asn Lys Ile Thr Leu Leu Glu Leu Glu 595 Asp Val Pro Leu Gln Pro Gln Val Gln Phe Leu Asp Lys Pro Ile Leu Asn Ser Thr Ser Thr Leu His Arg Thr His Ile Asn Ser Leu Ser Ala 625 Asp Thr Leu Ser Ala Ser Glu Pro Met Glu Leu Ser Gly His 645 <210> 178 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 178 24 tggctactcc aagaccctgg catg <210> 179 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 179 24 tggacaaatc cccttgctca gccc <210> 180 <211> 50 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 180 50 gggcttcacc gaagcagtgg acctttattt tgaccacctg atgtccaggg <210> 181 <211> 22 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 181 22 ccagctatga ctatgatgca cc <210> 182 <211> 24

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Glu Leu Glu Arg Ile Pro His Ala Ile Phe Ser Leu Ser Asn Leu Gln 290 295 300

Glu Leu Asp Leu Lys Ser Asn Asn Ile Arg Thr Ile Glu Glu Ile Ile Page 93

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gcaacagagg tgaaactact gggccatgga cagccactta actggatttc tttggagcaa 1380
                                                                              Page 98
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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ttaaattacc catgtaacca tttaactct ccagtgcact ttgccattaa agtctcttca 1680
cattgatttg tttccatgtg tgactcagag gtgagaattt tttcacatta tagtagcaag 1740 gaattggtgg tattatggac cgaactgaaa attttatgtt gaagccatat cccccatgat 1800
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                                                                               2280
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                                                                               2362
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Leu Leu Leu Leu Pro Pro Pro Pro Cys Pro Ala His Ser Ala Thr
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Trp Phe Asp Gln Ala Lys Phe Gly Ile Phe Ile His Trp Gly Val Phe 50 60
Ser Val Pro Ser Phe Gly Ser Glu Trp Phe Trp Trp Tyr Trp Gln Lys
65 70 75 80
Glu Lys Ile Pro Lys Tyr Val Glu Phe Met Lys Asp Asn Tyr Pro Pro 85 90 95
Ser Phe Lys Tyr Glu Asp Phe Gly Pro Leu Phe Thr Ala Lys Phe Phe 100 \hspace{1cm} 105 \hspace{1cm} 110
Asn Ala Asn Gln Trp Ala Asp Ile Phe Gln Ala Ser Gly Ala Lys Tyr
115 120 . 125
Ile Val Leu Thr Ser Lys His His Glu Gly Phe Thr Leu Trp Gly Ser
130 135 140
Glu Tyr Ser Trp Asn Trp Asn Ala Ile Asp Glu Gly Pro Lys Arg Asp
145 150 155 160
Ile Val Lys Glu Leu Glu Val Ala Ile Arg Asn Arg Thr Asp Leu Arg
165 170 175
Phe Gly Leu Tyr Tyr Ser Leu Phe Glu Trp Phe His Pro Leu Phe Leu
180 185 190
Glu Asp Glu Ser Ser Phe His Lys Arg Gln Phe Pro Val Ser Lys
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
Thr Leu Pro Glu Leu Tyr Glu Leu Val Asn Asn Tyr Gln Pro Glu Val
210 215 220
Leu Trp Ser Asp Gly Asp Gly Gly Ala Pro Asp Gln Tyr Trp Asn Ser
225 230 235 240
Thr Gly Phe Leu Ala Trp Leu Tyr Asn Glu Ser Pro Val Arg Gly Thr
245 250 255
val val Thr Asn Asp Arg Trp Gly Ala Gly Ser Ile Cys Lys His Gly 260 265 270
Gly Phe Tyr Thr Cys Ser Asp Arg Tyr Asn Pro Gly His Leu Leu Pro 275 280 285
His Lys Trp Glu Asn Cys Met Thr Ile Asp Lys Leu Ser Trp Gly Tyr
290 295 300
Arg Arg Glu Ala Gly Ile Ser Asp Tyr Leu Thr Ile Glu Glu Leu Val
305 310 315 320
Lys Gln Leu Val Glu Thr Val Ser Cys Gly Gly Asn Leu Leu Met Asn 325 330 335
Ile Gly Pro Thr Leu Asp Gly Thr Ile Ser Val Val Phe Glu Glu Arg
340 345 350
Leu Arg Gln Val Gly Ser Trp Leu Lys Val Asn Gly Glu Ala Ile Tyr
355 360 365
Glu Thr Tyr Thr Trp Arg Ser Gln Asn Asp Thr Val Thr Pro Asp Val
370 375 380
Trp Tyr Thr Ser Lys Pro Lys Glu Lys Leu Val Tyr Ala Ile Phe Leu 385 390 395 400
Lys Trp Pro Thr Ser Gly Gln Leu Phe Leu Gly His Pro Lys Ala Ile
405 410 415
Leu Gly Ala Thr Glu Val Lys Leu Leu Gly His Gly Gln Pro Leu Asn 420 430
              420
Trp Ile Ser Leu Glu Gln Asn Gly Ile Met Val Glu Leu Pro Gln Leu 435 440 445
Thr Ile His Gln Met Pro Cys Lys Trp Gly Trp Ala Leu Ala Leu Thr
450 455 460
Asn Val Ile
465
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<400> 196
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 197
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<210> 198
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
        oligonucleotide probe
<400> 198
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<210> 199
<211> 45
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
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                                                                                           .45
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<210> 200
<211> 2372
<212> DNA
<213> Homo sapiens
<400> 200
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gatcccatat tcaacactca aactgcaaca caaacaacag aatttattgt cagtgacagt 660
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tgaggagaca cacctgaggc tggtttcttt catgctcctt accctgcccc agctggggaa 1200
atcăaaaggg ccaaagaacc aaagaagaaa gtccaccctt ggttcctaac tggaatcagc 1260
                                                    Page 101
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1,txt
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gtcctaataa tatcccactg ggagaaagga gttttgcaaa gtgcaaggac ctaaaacatc 1440
tcatcagtat ccagtggtaa aaaggcctcc tggctgtctg aggctaggtg ggttgaaagc 1500
caaggagtca ctgagaccaa ggctttctct actgattccg cagctcagac cctttcttca 1560
gctctgaaag agaaacacgt atcccacctg acatgtcctt ctgagcccgg taagagcaaa 1620
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<211> 322
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<213> Artificial sequence
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<223> Synthetic protein
<400> 201
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35 40 45
Asn Gln Gln Leu Asn Phe Thr Glu Ala Lys Glu Ala Cys Arg Leu Leu 50 60
Gly Leu Ser Leu Ala Gly Lys Asp Gln Val Glu Thr Ala Leu Lys Ala 65 70. 75 80
Ser Phe Glu Thr Cys Ser Tyr Gly Trp Val Gly Asp Gly Phe Val Val
85 90 95
Ile Ser Arg Ile Ser Pro Asn Pro Lys Cys Gly Lys Asn Gly Val Gly 100 105 110
Val Leu Ile Trp Lys Val Pro Val Ser Arg Gln Phe Ala Ala Tyr Cys
Tyr Asn Ser Ser Asp Thr Trp Thr Asn Ser Cys Ile Pro Glu Ile Ile
130 135 140
Thr Thr Lys Asp Pro Ile Phe Asn Thr Gln Thr Ala Thr Gln Thr Thr
                                                                         160
Glu Phe Ile Val Ser Asp Ser Thr Tyr Ser Val Ala Ser Pro Tyr
165 170 175
Thr Ile Pro Ala Pro Thr Thr Pro Pro Ala Pro Ala Ser Thr Ser
                                              Page 102
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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            180
Ile Pro Arg Arg Lys Lys Leu Ile Cys Val Thr Glu Val Phe Met Glu
                              200
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Ala Phe Lys Asn Glu Ala Ala Gly Phe Gly Gly Val Pro Thr Ala Leu
225 230 235 240
Leu Val Leu Ala Leu Leu Phe Phe Gly Ala Ala Ala Gly Leu Gly Phe 245 250 255
Cys Tyr Val Lys Arg Tyr Val Lys Ala Phe Pro Phe Thr Asn Lys Asn 260 270
Gln Gln Lys Glu Met Ile Glu Thr Lys Val Val Lys Glu Glu Lys Ala
Asn Asp Ser Asn Pro Asn Glu Glu Ser Lys Lys Thr Asp Lys Asn Pro 290 295
Glu Glu Ser Lys Ser Pro Ser Lys Thr Thr Val Arg Cys Leu Glu Ala
                                           315
                     310
305
Glu Val
<210> 202
<211> 24
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<223> Description of Artificial Sequence: Synthetic
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<400> 202
                                                                        24
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<210> 203
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<400> 203
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<210> 204
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      oligonucleotide probe
<400> 204
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<223> Description of Artificial Sequence: Synthetic
           oligonucleotide probe
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<210> 206
<211> 1620
<212> DNA
<213> Homo sapiens
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<223> a, t, c or g
<220>
<221> modified_base
<222> (977)..(977)
<223> a, t, c or g
<220>
<221> modified_base
<222> (996)..(996)
<223> a, t, c or g
<220>
<221> modified_base
<222> (1003)..(1003)
<223> a, t, c or g
<400> 206
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                                                                         Page 104
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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actttccttt gtgtggtagg acttggagga gaaatccct ggactttcac taaccctctg 1560 acatactcc cacacccagt tgatggcttt ccgtaataaa aagattggga tttccttttg 1620
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<211> 296
<212> PRT
<213> Homo sapiens
<400> 207
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Ser Ala Ala Phe Leu Leu Val Arg Lys Leu Pro Pro Leu Cys His Gly 35 40 45
Leu Pro Thr Gln Arg Glu Asp Gly Asn Pro Cys Asp Phe Asp Trp Arg 50 55 60
Glu Val Glu Ile Leu Met Phe Leu Ser Ala Ile Val Met Met Lys Asn
65 70 75 80
Arg Arg Ser Ile Thr Val Glu Gln His Ile Gly Asn Ile Phe Met Phe 85 90 95
Ser Lys Val Ala Asn Thr Ile Leu Phe Phe Arg Leu Asp Ile Arg Met 100 \hspace{1cm} 105 \hspace{1cm} 110
Gly Leu Leu Tyr Ile Thr Leu Cys Ile Val Phe Leu Met Thr Cys Lys
115 120 125
Pro Pro Leu Tyr Met Gly Pro Glu Tyr Ile Lys Tyr Phe Asn Asp Lys
130 135 140
Thr Ile Asp Glu Glu Leu Glu Arg Asp Lys Arg Val Thr Trp Ile Val
145 150 155 160
Glu Phe Phe Ala Asn Trp Ser Asn Asp Cys Gln Ser Phe Ala Pro Ile
165 170 175
Tyr Ala Asp Leu Ser Leu Lys Tyr Asn Cys Thr Gly Leu Asn Phe Gly 180 185 190
Lys Val Asp Val Gly Arg Tyr Thr Asp Val Ser Thr Arg Tyr Lys Val
195 200 205
Ser Thr Ser Pro Leu Thr Lys Gln Leu Pro Thr Leu Ile Leu Phe Gln 210 220
Gly Gly Lys Glu Ala Met Arg Arg Pro Gln Ile Asp Lys Lys Gly Arg
225 230 235 240
Ala Val Ser Trp Thr Phe Ser Glu Glu Asn Val Ile Arg Glu Phe Asn
Leu Asn Glu Leu Tyr Gln Arg Ala Lys Lys Leu Ser Lys Ala Gly Asp
260 265 270
Asn Ile Pro Glu Glu Gln Pro Val Ala Ser Thr Pro Thr Thr Val Ser
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<211> 24
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       oligonucleotide probe
<400> 208
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<210> 209
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 209
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<210> 210
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
       oligonucleotide probe
<400> 210
                                                                               24
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<211> 50
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<212> DNA
<213> Homo sapiens
<400> 212
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                                              Page 106
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Gln Asn Leu Gly Asn Gly His Gly Lys Asp Leu Leu Asn Gly Val Lys
35 40 45
Leu Val Val Glu Thr Pro Glu Glu Thr Leu Phe Thr Tyr Gln Gly Ala 50 60
Ser Val Ile Leu Pro Cys Arg Tyr Arg Tyr Glu Pro Ala Leu Val Ser
65 70 75 80
Pro Arg Arg Val Arg Val Lys Trp Trp Lys Leu Ser Glu Asn Gly Ala
85 90 95
Pro Glu Lys Asp Val Leu Val Ala Ile Gly Leu Arg His Arg Ser Phe 100 105 110
Gly Asp Tyr Gln Gly Arg Val His Leu Arg Gln Asp Lys Glu His Asp
115 120 125
Val Ser Leu Glu Ile Gln Asp Leu Arg Leu Glu Asp Tyr Gly Arg Tyr
130 135 140
                                                          Page 107
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Tyr Gln Phe Asn Phe His Glu Gly Gln Gln Val Cys Ala Glu Gln Ala
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Ala Val Val Ala Ser Phe Glu Gln Leu Phe Arg Ala Trp Glu Glu Gly 195 200 205
Leu Asp Trp Cys Asn Ala Gly Trp Leu Gln Asp Ala Thr Val Gln Tyr 210 215 220
Pro Ile Met Leu Pro Arg Gln Pro Cys Gly Gly Pro Gly Leu Ala Pro 225 230 235 240
Gly Val Arg Ser Tyr Gly Pro Arg His Arg Arg Leu His Arg Tyr Asp
245 250 255
Val Phe Cys Phe Ala Thr Ala Leu Lys Gly Arg Val Tyr Tyr Leu Glu
260 265 270
His Pro Glu Lys Leu Thr Leu Thr Glu Ala Arg Glu Ala Cys Gln Glu
275 280 285
Asp Asp Ala Thr Ile Ala Lys Val Gly Gln Leu Phe Ala Ala Trp Lys 290 295 300
Phe His Gly Leu Asp Arg Cys Asp Ala Gly Trp Leu Ala Asp Gly Ser 305 310 315 320
Val Arg Tyr Pro Val Val His Pro His Pro Asn Cys Gly Pro Pro Glu
325 330 335
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355 360
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cgacgggcgt gtgggttggg aggagctgcg caacgccacc tatggccact acgcgcccgg 480
tgaagaattt catgacgtgg aggatgcaga gacctacaaa aagatgctgg ctcgggacga 540
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                                                                            780
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His Gly Asn Phe Gln Tyr Asp His Glu Ala Phe Leu Gly Arg Glu Val 50 55 60
Ala Lys Glu Phe Asp Gln Leu Thr Pro Glu Glu Ser Gln Ala Arg Leu
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Gly Arg Ile Val Asp Arg Met Asp Arg Ala Gly Asp Gly Asp Gly Trp
85 90 95
Val Ser Leu Ala Glu Leu Arg Ala Trp Ile Ala His Thr Gln Gln Arg
100 105. 110
His Ile Arg Asp Ser Val Ser Ala Ala Trp Asp Thr Tyr Asp Thr Asp
115 120 125
Arg Asp Gly Arg Val Gly Trp Glu Glu Leu Arg Asn Ala Thr Tyr Gly 130 140
His Tyr Ala Pro Gly Glu Glu Phe His Asp Val Glu Asp Ala Glu Thr
145 150 155 160
Tyr Lys Lys Met Leu Ala Arg Asp Glu Arg Arg Phe Arg Val Ala Asp
165 170 175
Gln Asp Gly Asp Ser Met Ala Thr Arg Glu Glu Leu Thr Ala Phe Leu
180 185 190
                                            Page 110
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Tyr Ile Ala Asp Leu Tyr Ser Ala Glu Pro Gly Glu Glu Glu Pro Ala
225 . 230 235 240
Trp Val Gln Thr Glu Arg Gln Gln Phe Arg Asp Phe Arg Asp Leu Asn 245 250 255
Lys Asp Gly His Leu Asp Gly Ser Glu Val Gly His Trp Val Leu Pro
260 265 270
Pro Ala Gln Asp Gln Pro Leu Val Glu Ala Asn His Leu Leu His Glu
Ser Asp Thr Asp Lys Asp Gly Arg Leu Ser Lys Ala Glu Ile Leu Gly 290 295 300
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 <211> 2403
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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         355
Ala Phe Leu Ile Asp Gly Ser Ser Ser Val Gly Asp Ser Asn Phe Arg 370 375 380
Leu Met Leu Glu Phe Val Ser Asn Ile Ala Lys Thr Phe Glu Ile Ser 385 390 395 400
385
Asp Ile Gly Ala Lys Ile Ala Ala Val Gln Phe Thr Tyr Asp Gln Arg 405 410 415
Thr Glu Phe Ser Phe Thr Asp Tyr Ser Thr Lys Glu Asn Val Leu Ala
420 425 430
Val Ile Arg Asn Ile Arg Tyr Met Ser Gly Gly Thr Ala Thr Gly Asp
435 440 445
Ala Ile Ser Phe Thr Val Arg Asn Val Phe Gly Pro Ile Arg Glu Ser
450 455 460
Pro Asn Lys Asn Phe Leu Val Ile Val Thr Asp Gly Gln Ser Tyr 465 470 475
Asp Val Gln Gly Pro Ala Ala Ala Ala His Asp Ala Gly Ile Thr Ile
                                        490
Phe Ser Val Gly Val Ala Trp Ala Pro Leu Asp Asp Leu Lys Asp Met 500 510
Ala Ser Lys Pro Lys Glu Ser His Ala Phe Phe Thr Arg Glu Phe Thr
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Phe Leu Glu Ser Gln Gln
545
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<211> 350
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Arg Lys Asp Ala Tyr Met Phe Trp Trp Leu Tyr Tyr Ala Thr Asn 60

Ser Cys Lys Asn Phe Ser Glu Leu Pro Leu Val Met Trp Leu Gln 75

Gly Gly Pro Gly Gly Ser Ser Thr Gly Phe Gly Asn Phe Glu Glu 90

Ile Gly Pro Leu Asp Ser Asp Leu Lys Pro Arg Lys Thr Thr Trp 105

Leu Gln Ala Ala Ser Leu Leu Phe Val Asp Asn Pro Val Gly Thr 120

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95 100 105
Met Pro Ser Phe Trp Ser Leu Gln Ala Tyr Tyr Thr Arg Tyr Phe 110 115 120
Val Ser Asn Ile Tyr Leu Ser Pro Arg Tyr Leu Gly Asn Ser Pro
125 130 135
Tyr Asp Ile Ala Leu Val Lys Leu Ser Ala Pro Val Thr Tyr Thr
140 145 150
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Glu Asn Arg Thr Asp Cys Trp Val Thr Gly Trp Gly Tyr Ile Lys
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Tyr Ser Phe Arg Lys Asp Ile Phe Gly Asp Met Val Cys Ala Gly 215 220 225
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His Gly Ala Gly Leu Phe Asp Val Thr Arg Gly Cys His Glu Ser 525

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^{212&}gt; PRT

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Arg Ser Pro Glu Asn Ser Leu Asp Pro Lys Met Ser Leu Thr Gly
Ala Arg Glu Leu Leu Pro His Ala Gln Glu Glu Ala Glu 350 355 360
Ala Glu Leu Pro Pro Ser Ser Glu Val Leu Ala Ser Val Phe Pro
                                                           375
Ala Gln Asp Lys Pro Gly Glu Leu Gln Ala Thr Leu Asp His Thr
                                      385
Gly His Thr Ser Ser Lys Ser Leu Pro Asn Phe Pro Asn Thr Ser
                                                           405
Ala Thr Ala Asn Ala Thr Gly Gly Arg Ala Leu Ala Leu Gln Ser
Ser Leu Pro Gly Ala Glu Gly Pro Asp Lys Pro Ser Val Val Ser
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Asn Asn Asn Glu Leu Glu Thr Ile Pro Asn Leu Gly Pro Val Ser
Ala Asn Ile Thr Leu Leu Ser Leu Ala Gly Asn Arg Ile Val Glu
Ile Leu Pro Glu His Leu Lys Glu Phe Gln Ser Leu Glu Thr Leu
Asp Leu Ser Ser Asn Asn Ile Ser Glu Leu Gln Thr Ala Phe Pro
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110 115 120
                                                          120
Ser Met Glu Pro Gly Tyr Phe Asp Asn Leu Ala Asn Thr Leu Leu
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt Ile Lys Gly Ser Asn Leu Ser Phe Ile Cys Ser Ala Ala Ser Ser 455 460 465 Ser Asp Ser Pro Met Thr Phe Ala Trp Lys Lys Asp Asn Glu Leu 480 Leu His Asp Ala Glu Met Glu Asn Tyr Ala His Leu Arg Ala Gln 495 485 Gly Gly Glu Val Met Glu Tyr Thr Thr Ile Leu Arg Leu Arg Glu 505 Val Glu Phe Ala Ser Glu Gly Lys Tyr Gln Cys Val Ile Ser Asn 515 520 525 His Phe Gly Ser Ser Tyr Ser Val Lys Ala Lys Leu Thr Val Asn 540 Met Leu Pro Ser Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 560 565 570 Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 575 580 585 Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val
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50 55 60 Ile Cys Val Arg Lys Asn Leu Arg Glu Val Pro Asp Gly Ile Ser
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt 290 Asn Pro Trp Asn Cys Asn Cys Asp Ile Leu Trp Leu Ser Trp Trp Pro Ser Asn Thr Ala Cys Cys Ala Arg 325 Ile Lys Asp Met Ala Asn Thr Pro Pro Asn Leu Lys Gly Arg Tyr Ile Gly Glu Leu Asp Gln Asn Tyr Phe Thr Cys Tyr Ala Pro Val Ile Val Glu Pro Pro Ala Asp Leu Asn Val Thr Glu Gly Met Ala Ala Glu Leu Lys Cys Arg Ala Ser Thr Ser Leu Thr Ser Val Ser Trp Ile Thr Pro Asn 380 385 Gly Thr Val Met Thr His Gly Ala Tyr Lys Val Arg Ile Ala Val 395 400 405 395 Leu Ser Asp Gly Thr Leu Asn Phe Thr Asn Val Thr Val Gin Asp Thr Gly Met Tyr Thr Cys Met Val Ser Asn Ser Val Gly Asn Thr 435 Thr Ala Ser Ala Thr Leu Asn Val Thr Ala Ala Thr Thr Thr Pro 440 Phe Ser Tyr Phe Ser Thr Val Thr Val Glu Thr Met Glu Pro Ser Gln Asp Glu Ala Arg Thr Thr Asp Asn Asn Val Gly Pro Thr Pro 470 475 480 480 val val Asp Trp Glu Thr Thr Asn val Thr Thr Ser Leu Thr Pro 495 Gln Ser Thr Arg Ser Thr Glu Lys Thr Phe Thr Ile Pro Val Thr 510 Asp Ile Asn Ser Gly Ile Pro Gly Ile Asp Glu Val Met Lys 515 520 Thr Lys Ile Ile Ile Gly Cys Phe Val Ala Ile Thr Leu Met Ala Ala Val Met Leu Val Ile Phe Tyr Lys Met Arg Lys Gln His His 550 Arg Gln Asn His His Ala Pro Thr Arg Thr Val Glu Ile Ile Asn 565 Val Asp Asp Glu Ile Thr Gly Asp Thr Pro Met Glu Ser His Leu 580 585 Pro Met Pro Ala Ile Glu His Glu His Leu Asn His Tyr Asn Ser Tyr Lys Ser Pro Phe Asn His Thr Thr Thr Val Asn Thr Ile Asn Page 158

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt 605 615

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545 550 555 Gly Gly Glu Val Met Glu Tyr Thr Thr Ile Leu Arg Leu Arg Val Glu Phe Ala Ser Glu Gly Lys Tyr Gln Cys Val Ile Ser Asn 575 580 585 His Phe Gly Ser Ser Tyr Ser Val Lys Ala Lys Leu Thr Val Asn 590 595 600 600 Met Leu Pro Ser Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg 615 Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 630 Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val Page 163

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705 Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 710 715 720 Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 725 730 735 Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln
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                                      1060
                1055
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50 55 60 Gln Pro Arg Cys Lys His Gly Glu Cys Ile Gly Pro Asn Lys Cys
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75 Lys Cys His Pro Gly Tyr Ala Gly Lys Thr Cys Asn Gln Asp Leu 80 85 90 Asn Glu Cys Gly Leu Lys Pro Arg Pro Cys Lys His Arg Cys Met 105 100 Asn Thr Tyr Gly Ser Tyr Lys Cys Tyr Cys Leu Asn Gly Tyr Met 110 115 120 Leu Met Pro Asp Gly Ser Cys Ser Ser Ala Leu Thr Cys Ser Met 125 130 135 Ala Asn Cys Gln Tyr Gly Cys Asp Val Val Lys Gly Gln Ile Arg 140 145 150 Cys Gln Cys Pro Ser Pro Gly Leu His Leu Ala Pro Asp Gly Arg 155 160 165 Thr Cys Val Asp Val Asp Glu Cys Ala Thr Gly Arg Ala Ser Cys 170 175 180 Pro Arg Phe Arg Gln Cys Val Asn Thr Phe Gly Ser Tyr Ile Cys 185 190 195 Lys Cys His Lys Gly Phe Asp Leu Met Tyr Ile Gly Gly Lys Tyr $200 \hspace{1cm} 205 \hspace{1cm} 210 \hspace{1cm}$ Gln Cys His Asp Ile Asp Glu Cys Ser Leu Gly Gln Tyr Gln Cys 225 Ser Ser Phe Ala Arg Cys Tyr Asn Val Arg Gly Ser Tyr Lys Cys 230 235 240 Lys Cys Lys Glu Gly Tyr Gln Gly Asp Gly Leu Thr Cys Val Tyr 245 250 255 Ile Pro Lys Val Met Ile Glu Pro Ser Gly Pro Ile His Val Pro 260 265 270 Lys Gly Asn Gly Thr Ile Leu Lys Gly Asp Thr Gly Asn Asn Asn 275 280 285 Trp Ile Pro Asp Val Gly Ser Thr Trp Trp Pro Pro Lys Thr Pro 290 295 300 Tyr Ile Pro Pro Ile Ile Thr Asn Arg Pro Thr Ser Lys Pro Thr Page 173

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                                       370
                                                            375
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                                                            390
                 380
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Lys Asp Asn Asp Leu His Trp Glu Pro Ile Arg Asp Pro Ala Gly
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                                       415
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425 430 430
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Ala Ala Arg Leu Val Leu Pro Leu Gly Arg Leu Met His Ser Gly
440 445 450
Asp Leu Cys Leu Ser Phe Arg His Lys Val Thr Gly Leu His Ser
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Gly Thr Leu Gln Val Phe Val Arg Lys His Gly Ala His Gly Ala
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His Thr Pro Ala Ser Asp Ile Gln Ile Ile Trp Leu Phe Glu Arg
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val val Pro Asp Leu Glu Tyr Gln His Lys Phe Thr Met Met Pro

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Glu Asp Ala Leu Asp Asp Phe Gly Ile Tyr Glu Phe Val Ala Phe

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35 40 45 Asp Trp Met Ile Glu Phe Tyr Ala Pro Trp Cys Pro Ala Cys Gln
50 55 60 Asn Leu Gln Pro Glu Trp Glu Ser Phe Ala Glu Trp Gly Glu Asp
65 70 75 Leu Glu Val Asn Ile Ala Lys Val Asp Val Thr Glu Gln Pro Gly 80 85 90 Leu Ser Gly Arg Phe Ile Ile Thr Ala Leu Pro Thr Ile Tyr His 95 100 105 Cys Lys Asp Gly Glu Phe Arg Arg Tyr Gln Gly Pro Arg Thr 110 115 Lys Asp Phe Ile Asn Phe Ile Ser Asp Lys Glu Trp Lys Ser Ile 125 130 135 Glu Pro Val Ser Ser Trp Phe Gly Pro Gly Ser Val Leu Met Ser 140 . 145 Ser Met Ser Ala Leu Phe Gln Leu Ser Met Trp Ile Arg Thr 160 His Asn Tyr Phe Ile Glu Asp Leu Gly Leu Pro Val Trp Gly Ser 175 180 Page 180

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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Ser Glu Ser Ala Gln Pro Leu Lys Lys Val Glu Glu Glu Gln Glu 240

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40
45
Glu Tyr Ile Leu Val Glu Glu Ala Lys Leu Ser Lys Ile Lys Ser
50 55 60
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170 175 180 **180** Asp Tyr Tyr His Thr Val Leu Trp Met Glu Gln Val Leu Lys Gln 185 190 195 Leu Asp Ala Gly Glu Glu Ala Thr Thr Lys Ser Gln Val Leu 200 205 210 Asp Tyr Leu Ser Tyr Ala Val Phe Gln Leu Gly Asp Leu His Arg 215 220 225 Ala Leu Glu Leu Thr Arg Arg Leu Leu Ser Leu Asp Pro Ser His 230 235 240 Glu Arg Ala Gly Gly Asn Leu Arg Tyr Phe Glu Gln Leu Leu Glu 245 250 255 Glu Glu Arg Glu Lys Thr Leu Thr Asn Gln Thr Glu Ala Glu Leu 260 265 270 Ala Thr Pro Glu Gly Ile Tyr Glu Arg Pro Val Asp Tyr Leu Pro 275 280 285 Glu Arg Asp Val Tyr Glu Ser Leu Cys Arg Gly Glu Gly Val Leu Thr Pro Arg Arg Gln Lys Arg Leu Phe Cys Arg Tyr His 305 Gly Asn Arg Ala Pro Gln Leu Leu Ile Ala Pro Phe Lys Glu Glu 320 325 330 Asp Glu Trp Asp Ser Pro His Ile Val Arg Tyr Tyr Asp Val Met 335 340 345 Ser Asp Glu Glu Ile Glu Arg Ile Lys Glu Ile Ala Lys Pro Lys 350 355 360 Leu Ala Arg Ala Thr Val Arg Asp Pro Lys Thr Gly Val Leu Thr Page 184

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Arg Pro Phe Asp Ser Gly Leu Lys Thr Glu Gly Asn Arg Leu Ala
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                                       460
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485 490 495
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50 55 60 Leu Asp Gln Ser Asp Glu Asp Phe Lys Pro Arg Ile Val Pro Tyr
65 70 75 Tyr Arg Asp Pro Asn Lys Pro Tyr Lys Lys Val Leu Arg Thr Arg 80 85 90 Tyr Ile Gln Thr Glu Leu Gly Ser Arg Glu Arg Leu Leu Val Ala 95 100 105 Val Leu Thr Ser Arg Ala Thr Leu Ser Thr Leu Ala Val Ala Val 110 115 Asn Arg Thr Val Ala His His Phe Pro Arg Leu Leu Tyr Phe Thr 125 130 135 Gly Gln Arg Gly Ala Arg Ala Pro Ala Gly Met Gln Val Val Ser 140 145 150 His Gly Asp Glu Arg Pro Ala Trp Leu Met Ser Glu Thr Leu Arg 155 160 165 His Leu His Thr His Phe Gly Ala Asp Tyr Asp Trp Phe Phe Ile 170 175 180 Met Gln Asp Asp Thr Tyr Val Gln Ala Pro Arg Leu Ala Ala Leu

Ala Gly His Leu Ser Ile Asn Gln Asp Leu Tyr Leu Gly Arg Ala 200 205 210 Glu Glu Phe Ile Gly Ala Gly Glu Gln Ala Arg Tyr Cys His Gly 215 220 225 Gly Phe Gly Tyr Leu Leu Ser Arg Ser Leu Leu Leu Arg Leu Arg 230 235 240 Pro His Leu Asp Gly Cys Arg Gly Asp Ile Leu Ser Ala Arg Pro 245 250 255 Asp Glu Trp Leu Gly Arg Cys Leu Ile Asp Ser Leu Gly Val 260 265 Cys Val Ser Gln His Gln Gly Gln Gln Tyr Arg Ser Phe Glu Leu 275 280 285 Ala Lys Asn Arg Asp Pro Glu Lys Glu Gly Ser Ser Ala Phe Leu 290 295 300 Ser Ala Phe Ala Val His Pro Val Ser Glu Gly Thr Leu Met Tyr Arg Leu His Lys Arg Phe Ser Ala Leu Glu Leu Glu Arg Ala Tyr 320 325 330 Ser Glu Ile Glu Gln Leu Gln Ala Gln Ile Arg Asn Leu Thr Val 335 340 345 Leu Thr Pro Glu Gly Glu Ala Gly Leu Ser Trp Pro Val Gly Leu 350 355 360 Pro Ala Pro Phe Thr Pro His Ser Arg Phe Glu Val Leu Gly 365 Asp Tyr Phe Thr Glu Gln His Thr Phe Ser Cys Ala Asp Gly Ala 380 385 390 380 Pro Lys Cys Pro Leu Gln Gly Ala Ser Arg Ala Asp Val Gly Asp 395 400 405 Ala Leu Glu Thr Ala Leu Glu Gln Leu Asn Arg Arg Tyr Gln Pro
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt tgtccccaag ccgttctaga cgcgggaaaa atgctttctg aaagcagctc 100 ctttttgaag ggtgtgatgc ttggaagcat tttctgtgct ttgatcacta 150 tgctaggaca cattaggatt ggtcatggaa atagaatgca ccaccatgag 200 catcatcacc tacaagctcc taacaaagaa gatatcttga aaatttcaga 250 ggatgagcgc atggagctca gtaagagctt tcgagtatac tgtattatcc 300 ttgtaaaacc caaagatgtg agtctttggg ctgcagtaaa ggagacttgg 350 accaaacact gtgacaaagc agagttcttc agttctgaaa atgttaaagt 400 gtttgagtca attaatatgg acacaaatga catgtggtta atgatgagaa 450. aagcttacaa atacgccttt gataagtata gagaccaata caactggttc 500 ttccttgcac gccccactac gtttgctatc attgaaaacc taaagtattt 550 tttgttaaaa aaggatccat cacagccttt ctatctaggc cacactataa 600 aatctggaga ccttgaatat gtgggtatgg aaggaggaat tgtcttaagt 650 gtagaatcaa tgaaaagact taacagcctt ctcaatatcc cagaaaagtg 700 tcctgaacag ggagggatga tttggaagat atctgaagat aaacagctag 750 cagtttgcct gaaatatgct ggagtatttg cagaaaatgc agaagatgct 800 gatggaaaag atgtatttaa taccaaatct gttgggcttt ctattaaaga 850 ggcaatgact tatcacccca accaggtagt agaaggctgt tgttcagata 900 tggctgttac ttttaatgga ctgactccaa atcagatgca tgtgatgatg 950 tatggggtat accgccttag ggcatttggg catattttca atgatgcatt 1000 ggttttctta cctccaaatg gttctgacaa tgactgagaa gtggtagaaa 1050 agcgtgaata tgatctttgt ataggacgtg tgttgtcatt atttgtagta 1100 gtaactacat atccaataca gctgtatgtt tctttttctt ttctaatttg 1150 gtggcactgg tataaccaca cattaaagtc agtagtacat ttttaaatga 1200 gggtggtttt tttctttaaa acacatgaac attgtaaatg tgttggaaag 1250 aagtgtttta agaataataa ttttgcaaat aaactattaa taaatattat 1300 atgtgataaa ttctaaatta tgaacattag aaatctgtgg ggcacatatt 1350 tttgctgatt ggttaaaaaa ttttaacagg tctttagcgt tctaagatat 1400 gcaaatgata tctctagttg tgaatttgtg attaaagtaa aacttttagc 1450 tgtgtgttcc ctttacttct aatactgatt tatgttctaa gcctccccaa 1500 gttccaatgg atttgccttc tcaaaatgta caactaagca actaaagaaa 1550 attaaagtga aagttgaaaa at 1572

<211> 318 <212> PRT

<213> Homo Sapien

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35 40 45 Ala Pro Asn Lys Glu Asp Ile Leu Lys Ile Ser Glu Asp Glu Arg
50 55 60 Met Glu Leu Ser Lys Ser Phe Arg Val Tyr Cys Ile Ile Leu Val Lys Pro Lys Asp Val Ser Leu Trp Ala Ala Val Lys Glu Thr Trp Thr Lys His Cys Asp Lys Ala Glu Phe Phe Ser Ser Glu Asn Val 95 100 105 Lys Val Phe Glu Ser Ile Asn Met Asp Thr Asn Asp Met Trp Leu 110 115 120 Met Met Arg Lys Ala Tyr Lys Tyr Ala Phe Asp Lys Tyr Arg Asp 125 130 135 Gln Tyr Asn Trp Phe Phe Leu Ala Arg Pro Thr Thr Phe Ala Ile Ile Glu Asn Leu Lys Tyr Phe Leu Leu Lys Lys Asp Pro Ser Gln. 165 Pro Phe Tyr Leu Gly His Thr Ile Lys Ser Gly Asp Leu Glu Tyr 170 175 180 Val Gly Met Glu Gly Gly Ile Val Leu Ser Val Glu Ser Met Lys 185 190 195 Arg Leu Asn Ser Leu Leu Asn Ile Pro Glu Lys Cys Pro Glu Gln
200 205 210 Gly Gly Met Ile Trp Lys Ile Ser Glu Asp Lys Gln Leu Ala Val 215 220 225 Cys Leu Lys Tyr Ala Gly Val Phe Ala Glu Asn Ála Glu Asp Ala 230 235 240 Asp Gly Lys Asp Val Phe Asn Thr Lys Ser Val Gly Leu Ser Ile 245 250 255 Lys Glu Ala Met Thr Tyr His Pro Asn Gln Val Val Glu Gly Cys Cys Ser Asp Met Ala Val Thr Phe Asn Gly Leu Thr Pro Asn Gln 275 280 285 Met His Val Met Met Tyr Gly Val Tyr Arg Leu Arg Ala Phe Gly Page 192

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt 295 His Ile Phe Asn Asp Ala Leu Val Phe Leu Pro Pro Asn Gly Ser 305 310 315 305 310 Asp Asn Asp <210> 342 <211> 23 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 342 tccccaagcc gttctagacg cgg 23 <210> 343 <211> 18 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 343 ctggttcttc cttgcacg 18 <210> 344 <211> 28 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 344 gcccaaatgc cctaaggcgg tatacccc 28 <210> 345 <211> 50 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 345 gggtgtgatg cttggaagca ttttctgtgc tttgatcact atgctaggac 50 <210> 346 <211> 25 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe

<400> 346

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<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 347
ccctcatgta ccggctcc 18
<210> 348
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 348
 ggattctaat acgactcact atagggctca gaaaagcgca acagagaa 48
<210> 349
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 349
 ctatgaaatt aaccctcact aaagggatgt cttccatgcc aaccttc 47
<210> 350
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
 ggattctaat acgactcact atagggcggc gatgtccact ggggctac 48
<210> 351
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 351
 ctatgaaatt aaccctcact aaagggacga ggaagatggg cggatggt 48
<210> 352
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<210> 353
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 353
ctatgaaatt aaccctcact aaagggacgg gggacaccac ggaccaga 48
<210> 354
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 354
 ggattctaat acgactcact atagggcttg ctgcggtttt tgttcctg 48
<210> 355
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 355
 ctatgaaatt aaccctcact aaagggagct gccgatccca ctggtatt 48
<210> 356
<211> 46
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 356
 ggattctaat acgactcact atagggcgga tcctggccgg cctctg 46
<210> 357
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 357
 ctatgaaatt aaccctcact aaagggagcc cgggcatggt ctcagtta 48
<210> 358
<211> 47
<212> DNA
<213> Artificial Sequence
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<220>
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<210> 359
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
 ctatgaaatt aaccctcact aaagggacca aggccacaaa cggaaatc 48
<210> 360
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 360
 ggattctaat acgactcact atagggctgt gctttcattc tgccagta 48
<210> 361
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 361
 ctatgaaatt aaccctcact aaagggaggg tacaattaag gggtggat 48
<210> 362
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 362
 ggattctaat acgactcact atagggcccg cctcgctcct gctcctg 47
<210> 363
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 363
ctatgaaatt aaccctcact aaagggagga ttgccgcgac cctcacag 48
<210> 364
<211> 47
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 364
ggattctaat acgactcact atagggcccc tcctgccttc cctgtcc 47
<210> 365
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 365
 ctatgaaatt aaccctcact aaagggagtg gtggccgcga ttatctgc 48
<210> 366
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 366
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<210> 367
<211> 47
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 367
 ctatgaaatt aaccctcact aaagggacag acggggcaga gggagtg 47
<210> 368
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 368
 ggattctaat acgactcact atagggccag gaggcgtgag gagaaac 47
<210> 369
<211> 48
<212> DNA
<213> Artificial Sequence
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<223> Synthetic Oligonucleotide Probe
<400> 369
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
ctatgaaatt aaccctcact aaagggaaag acatgtcatc gggagtgg 48
<210> 370
<211> 48
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 370
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<210> 371
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 371
 ctatgaaatt aaccctcact aaagggacac agacagagcc ccatacgc 48
<210> 372
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 372
 ggattctaat acgactcact atagggccag ggaaatccgg atgtctc 47
<210> 373
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 373
 ctatgaaatt aaccctcact aaagggagta aggggatgcc accgagta 48
<210> 374
<211> 47
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 374
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<210> 375 <211> 48 <212> DNA

<220>

<213> Artificial Sequence

CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt <223> Synthetic Oligonucleotide Probe <400> 375 ctatgaaatt aaccctcact aaagggatcc caggtgatga ggtccaga 48 <210> 376 <211> 997 <212> DNA <213> Homo Sapien <400> 376 cccacgcgtc cgatcttacc aacaaaacac tcctgaggag aaagaaagag 50 aaaaaatgaa ttcatctaaa tcatctgaaa cacaatgcac agagagagga 150 tgcttctctt cccaaatgtt cttatggact gttgctggga tccccatcct 200 atttctcagt gcctgtttca tcaccagatg tgttgtgaca tttcgcatct 250 ttcaaacctg tgatgagaaa aagtttcagc tacctgagaa tttcacagag 300 ctctcctgct acaattatgg atcaggttca gtcaagaatt gttgtccatt 350 gaactgggaa tattttcaat ccagctgcta cttcttttct actgacacca 400 tttcctgggc gttaagttta aagaactgct cagccatggg ggctcacctg 450 taaaatgaga gagttttta ttggactgtc agaccaggtt gtcgagggtc 550 agtggcaatg ggtggacggc acacctttga caaagtctct gagcttctgg 600 gatgtagggg agcccaacaa catagctacc ctggaggact gtgccaccat 650 gagagactct tcaaacccaa ggcaaaattg gaatgatgta acctgtttcc 700 tcaattattt tcggatttgt gaaatggtag gaataaatcc tttgaacaaa 750 ggaaaatctc tttaagaaca gaaggcacaa ctcaaatgtg taaagaagga 800 agagcaagaa catggccaca cccaccgccc cacacgagaa atttgtgcgc 850 tgaacttcaa aggacttcat aagtatttgt tactctgata caaataaaaa 900 <210> 377 <211> 219 <212> PRT <213> Homo Sapien <400> 377 Met Asn Ser Ser Lys Ser Ser Glu Thr Gln Cys Thr Glu Arg Gly Cys Phe Ser Ser Gln Met Phe Leu Trp Thr Val Ala Gly Ile Pro

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
Ile Leu Phe Leu Ser Ala Cys Phe Ile Thr Arg Cys Val Val Thr
Phe Arg Ile Phe Gln Thr Cys Asp Glu Lys Lys Phe Gln Leu Pro
50 55 60
Glu Asn Phe Thr Glu Leu Ser Cys Tyr Asn Tyr Gly Ser Gly Ser
65 70 75
val Lys Asn Cys Cys Pro Leu Asn Trp Glu Tyr Phe Gln Ser Ser
80 85 90
Cys Tyr Phe Phe Ser Thr Asp Thr Ile Ser Trp Ala Leu Ser Leu 95 100 105
Lys Asn Cys Ser Ala Met Gly Ala His Leu Val Val Ile Asn Ser
Gln Glu Glu Gln Glu Phe Leu Ser Tyr Lys Lys Pro Lys Met Arg
Glu Phe Phe Ile Gly Leu Ser Asp Gln Val Val Glu Gly Gln Trp
Gln Trp Val Asp Gly Thr Pro Leu Thr Lys Ser Leu Ser Phe Trp
155 160 165
Asp Val Gly Glu Pro Asn Asn Ile Ala Thr Leu Glu Asp Cys Ala
Thr Met Arg Asp Ser Ser Asn Pro Arg Gln Asn Trp Asn Asp Val
Thr Cys Phe Leu Asn Tyr Phe Arg Ile Cys Glu Met Val Gly Ile
                                                             210
                 200
Asn Pro Leu Asn Lys Gly Lys Ser Leu
215
<210> 378
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 378
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<210> 379
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 379
tattcctacc atttcacaaa tccg 24
<210> 380
<211> 49
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 380
 ggaggactgt gccaccatga gagactcttc aaacccaagg caaaattgg 49
<210> 381
<211> 26
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 381
 gcagattttg aggacagcca cctcca 26
<210> 382
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 382
 ggccttgcag acaaccgt 18
<210> 383
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 383
 cagactgagg gagatccgag a 21
<210> 384
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 384
 cagctgccct tccccaacca 20
<210> 385
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 385
 catcaagcgc ctctacca 18
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<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 386
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<210> 387
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 387
 gggccatcac agctccct 18
<210> 388
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 388
 gggatgtggt gaacacagaa ca 22
<210> 389
<211> 22
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<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 389
 tgccagctgc atgctgccag tt 22
<210> 390
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 390
cagaaggatg tcccgtggaa 20
<210> 391
<211> 17
<212> DNA
<213> Artificial Sequence
<220>
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<223> Synthetic oligonucleotide probe
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<210> 392
<211> 21
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<223> Synthetic oligonucleotide probe
<400> 392
 gacggcatcc tcagggccac a 21
<210> 393
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 393
 atgtcctcca tgcccacgcg 20
<210> 394
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 394
 gagtgcgaca tcgagagctt 20
<210> 395
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 395
 ccgcagcctc agtgatga 18
<210> 396
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 396
 gaagagcaca gctgcagatc c 21
<210> 397
<211> 22
<212> DNA
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<223> Synthetic oligonucleotide probe
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<210> 398
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 398
 cctctggcgc ccccactcaa 20
<210> 399
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 399
 ccaggagagc tggcgatg 18
<210> 400
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 400
 gcaaattcag ggctcactag aga 23
<210> 401
<211> 29
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 401
 cacagagcat ttgtccatca gcagttcag 29
<210> 402
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 402
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ggcagagact tccagtcact ga 22

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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
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<211> 22
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<223> Synthetic oligonucleotide probe
<400> 403
gccaagggtg gtgttagata gg 22
<210> 404
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 404
 caggccccct tgatctgtac ccca 24
<210> 405
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 405
 gggacgtgct tctacaagaa cag 23
<210> 406
<211> 26
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 406
 caggettaca atgttatgat cagaca 26
<210> 407
<211> 31
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 407
 tattcagagt tttccattgg cagtgccagt t 31
<210> 408
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<400> 408
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<210> 409
<211> 23
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<223> Synthetic oligonucleotide probe
<400> 409
 cgatcttctc cacccaggag cgg 23
<210> 410
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 410
 gccaggcctc acattcgt 18
<210> 411
<211> 23
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 411
 ctccctgaat ggcagcctga gca 23
<210> 412
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 412
 aggtgtttat taagggccta cgct 24
 <210> 413
 <211> 19
 <212> DNA
 <213> Artificial Sequence
 <223> Synthetic oligonucleotide probe
 <400> 413
 cagagcagag ggtgccttg 19
 <210> 414
 <211> 21
 <212> DNA
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<213> Artificial Sequence

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<223> Synthetic oligonucleotide probe
<400> 414
 tggcggagtc ccctcttggc t 21
<210> 415
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 415
ccctgtttcc ctatgcatca ct 22
<210> 416
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 416
 tcaaccctg accctttcct a 21
<210> 417
<211> 24
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 417
 ggcaggggac aagccatctc tcct 24
<210> 418
<211> 20
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<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 418
 gggactgaac tgccagcttc 20
<210> 419
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Synthetic oligonucleotide probe
<400> 419
 gggccctaac ctcattacct tt 22
<210> 420
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CORRECTED SEQUENCE LISTING FROM 10466.14 01.29.02P1618P2C1.txt
<211> 23
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<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 420
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<210> 421
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 421
 tctgtccacc atcttgcctt g 21
<210> 422
<211> 3554
<212> DNA
<213> Homo Sapien
<400> 422
 gggactacaa gccgcgccgc gctgccgctg gcccctcagc aaccctcgac 50
 atggcgctga ggcggccacc gcgactccgg ctctgcgctc ggctgcctga 100
 cttcttcctg ctgctgcttt tcaggggctg cctgataggg gctgtaaatc 150
 tcaaatccag caatcgaacc ccagtggtac aggaatttga aagtgtggaa 200
 ctgtcttgca tcattacgga ttcgcagaca agtgacccca ggatcgagtg 250
 gaagaaaatt caagatgaac aaaccacata tgtgtttttt gacaacaaaa 300
 ttcagggaga cttggcgggt cgtgcagaaa tactggggaa gacatccctg 350
 aagatctgga atgtgacacg gagagactca gccctttatc gctgtgaggt 400
 cgttgctcga aatgaccgca aggaaattga tgagattgtg atcgagttaa 450
 ctgtgcaagt gaagccagtg acccctgtct gtagagtgcc gaaggctgta 500
 ccagtaggca agatggcaac actgcactgc caggagagtg agggccaccc 550
 ccggcctcac tacagctggt atcgcaatga tgtaccactg cccacggatt 600
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